

FIG. 1

FIG. 2 (1 of 35)

Atom								
Type	Residue		X	Y	Z			
ATOM	1	CA MET	1	-24.866	15.635	-19.264	1.00	15.51
ATOM	2	HA MET	1	-25.390	14.593	-19.279	1.00	15.31
ATOM	3	CB MET	1	-25.849	16.777	-18.999	1.00	15.70
ATOM	4	HB1 MET	1	-26.446	16.943	-19.993	1.00	15.58
ATOM	5	HB2 MET	1	-25.298	17.678	-18.764	1.00	15.98
ATOM	6	CG MET	1	-26.765	16.422	-17.927	1.00	15.98
ATOM	7	HG1 MET	1	-26.172	16.103	-16.986	1.00	16.14
ATOM	8	HG2 MET	1	-27.432	15.624	-18.121	1.00	15.63
ATOM	9	SD MET	1	-27.737	17.976	-17.362	1.00	16.37
ATOM	10	CE MET	1	-28.779	17.948	-18.839	1.00	15.46
ATOM	11	HE1 MET	1	-28.342	17.340	-19.619	1.00	15.46
ATOM	12	HE2 MET	1	-28.851	18.969	-19.180	1.00	16.49
ATOM	13	HE3 MET	1	-29.765	17.590	-19.600	1.00	16.65
ATOM	14	C MET	1	-24.152	15.871	-20.591	1.00	15.23
ATOM	15	O MET	1	-23.015	16.304	-20.628	1.00	15.46
ATOM	16	N MET	1	-23.891	15.675	-18.134	1.00	16.16
ATOM	17	HT1 MET	1	-23.122	15.001	-18.319	1.00	16.50
ATOM	18	HT2 MET	1	-24.371	15.419	-17.249	1.00	16.33
ATOM	19	HT3 MET	1	-23.495	16.634	-18.051	1.00	16.24
ATOM	20	N ALA	2	-24.826	15.613	-21.680	1.00	14.89
ATOM	21	HN ALA	2	-25.749	15.283	-21.617	1.00	14.81
ATOM	22	CA ALA	2	-24.206	15.846	-23.013	1.00	14.80
ATOM	23	HA ALA	2	-23.256	15.339	-23.085	1.00	15.17
ATOM	24	CB ALA	2	-25.199	15.268	-24.026	1.00	15.06
ATOM	25	HB1 ALA	2	-25.968	15.998	-24.234	1.00	15.17
ATOM	26	HB2 ALA	2	-25.650	14.375	-23.618	1.00	15.16
ATOM	27	HB3 ALA	2	-24.680	15.021	-24.941	1.00	15.20
ATOM	28	C ALA	2	-24.035	17.355	-23.207	1.00	14.21
ATOM	29	O ALA	2	-24.023	18.097	-22.248	1.00	13.70
ATOM	30	N ALA	3	-23.920	17.811	-24.431	1.00	14.43
ATOM	31	HN ALA	3	-23.947	17.183	-25.183	1.00	14.93
ATOM	32	CA ALA	3	-23.761	19.283	-24.704	1.00	14.09
ATOM	33	HA ALA	3	-23.704	19.448	-25.769	1.00	14.05
ATOM	34	CB ALA	3	-25.032	19.954	-24.154	1.00	14.73
ATOM	35	HB1 ALA	3	-25.766	19.196	-23.921	1.00	15.01
ATOM	36	HB2 ALA	3	-25.435	20.624	-24.899	1.00	14.84
ATOM	37	HB3 ALA	3	-24.797	20.512	-23.260	1.00	14.95
ATOM	38	C ALA	3	-22.497	19.847	-24.031	1.00	13.47
ATOM	39	O ALA	3	-21.504	20.099	-24.689	1.00	13.42
ATOM	40	N GLY	4	-22.521	20.052	-22.737	1.00	13.16
ATOM	41	HN GLY	4	-23.126	19.842	-22.224	1.00	13.14
ATOM	42	CA GLY	4	-21.324	20.601	-22.038	1.00	12.72
ATOM	43	HA1 GLY	4	-21.640	21.265	-21.248	1.00	13.03
ATOM	44	HA2 GLY	4	-20.715	21.145	-22.745	1.00	12.88
ATOM	45	C GLY	4	-20.510	19.454	-21.438	1.00	11.83
ATOM	46	O GLY	4	-20.666	18.306	-21.810	1.00	11.80
ATOM	47	N GLN	5	-19.639	19.764	-20.517	1.00	11.27
ATOM	48	HN GLN	5	-19.534	20.700	-20.246	1.00	11.46
ATOM	49	CA GLN	5	-18.796	18.703	-19.882	1.00	10.54
ATOM	50	HA GLN	5	-18.209	18.192	-20.629	1.00	10.74
ATOM	51	CB GLN	5	-17.867	19.456	-18.918	1.00	10.66
ATOM	52	HB1 GLN	5	-17.255	20.149	-19.477	1.00	10.48
ATOM	53	HB2 GLN	5	-17.231	18.747	-18.408	1.00	10.80
ATOM	54	CG GLN	5	-18.696	20.230	-17.882	1.00	11.16
ATOM	55	HG1 GLN	5	-18.846	19.612	-17.008	1.00	11.36
ATOM	56	HG2 GLN	5	-19.655	20.493	-18.302	1.00	11.17
ATOM	57	CD GLN	5	-17.944	21.499	-17.481	1.00	11.68
ATOM	58	OE1 GLN	5	-17.092	21.465	-16.617	1.00	11.76
ATOM	59	NE2 GLN	5	-18.222	22.627	-18.079	1.00	12.23
ATOM	60	HE21 GLN	5	-18.910	22.655	-18.776	1.00	12.30
ATOM	61	HE22 GLN	5	-17.740	23.443	-17.933	1.00	12.69
ATOM	62	C GLN	5	-19.665	17.702	-19.111	1.00	9.75
ATOM	63	O GLN	5	-20.496	18.080	-18.306	1.00	9.62
ATOM	64	N ASN	6	-19.460	16.426	-19.339	1.00	9.41

ATOM	55	HN	ASN	6	-18.775	15.151	-19.981	1.00	9.70	SEG1
ATOM	56	CA	ASN	6	-20.257	15.396	-18.650	1.00	8.79	SEG1
ATOM	57	HA	ASN	6	-21.287	15.710	-18.535	1.00	9.04	SEG1
ATOM	63	CB	ASN	6	-20.161	14.093	-19.418	1.00	9.02	SEG1
ATOM	69	HB1	ASN	6	-20.676	14.225	-20.357	1.00	8.79	SEG1
ATOM	70	HB2	ASN	6	-20.631	13.293	-18.867	1.00	9.43	SEG1
ATOM	71	CG	ASN	6	-18.700	13.710	-19.695	1.00	9.37	SEG1
ATOM	72	OD1	ASN	6	-17.847	13.872	-18.345	1.00	9.72	SEG1
ATOM	73	ND2	ASN	6	-18.379	13.216	-20.863	1.00	9.56	SEG1
ATOM	74	HD21	ASN	6	-19.069	13.099	-21.547	1.00	9.44	SEG1
ATOM	75	HD22	ASN	6	-17.450	12.985	-21.056	1.00	9.96	SEG1
ATOM	76	C	ASN	6	-19.675	15.229	-17.193	1.00	8.00	SEG1
ATOM	77	O	ASN	6	-18.499	15.450	-16.975	1.00	7.88	SEG1
ATOM	78	N	GLY	7	-20.492	14.873	-16.233	1.00	7.69	SEG1
ATOM	79	HN	GLY	7	-21.441	14.724	-16.431	1.00	8.01	SEG1
ATOM	80	CA	GLY	7	-19.993	14.729	-14.831	1.00	7.15	SEG1
ATOM	81	HA1	GLY	7	-20.823	14.807	-14.147	1.00	7.30	SEG1
ATOM	82	HA2	GLY	7	-19.286	15.520	-14.625	1.00	7.43	SEG1
ATOM	83	C	GLY	7	-19.305	13.377	-14.636	1.00	6.36	SEG1
ATOM	84	O	GLY	7	-19.058	12.651	-15.579	1.00	6.40	SEG1
ATOM	85	N	HIS	8	-18.990	13.040	-13.407	1.00	5.93	SEG1
ATOM	86	HN	HIS	8	-19.199	13.649	-12.668	1.00	6.17	SEG1
ATOM	87	CA	HIS	8	-18.311	11.740	-13.127	1.00	5.39	SEG1
ATOM	88	HA	HIS	8	-18.677	10.972	-13.791	1.00	5.57	SEG1
ATOM	89	CB	HIS	8	-16.830	12.005	-13.398	1.00	5.92	SEG1
ATOM	90	HB1	HIS	8	-16.355	12.352	-12.492	1.00	6.12	SEG1
ATOM	91	HB2	HIS	8	-16.733	12.757	-14.166	1.00	6.23	SEG1
ATOM	92	CG	HIS	8	-16.170	10.733	-13.851	1.00	6.32	SEG1
ATOM	93	ND1	HIS	8	-16.762	9.885	-14.773	1.00	6.76	SEG1
ATOM	94	HD1	HIS	8	-17.627	10.024	-15.215	1.00	6.84	SEG1
ATOM	95	CD2	HIS	8	-14.976	10.147	-13.515	1.00	6.77	SEG1
ATOM	96	HD2	HIS	8	-14.259	10.552	-12.816	1.00	6.87	SEG1
ATOM	97	CE1	HIS	8	-15.933	8.842	-14.956	1.00	7.39	SEG1
ATOM	98	HE1	HIS	8	-16.140	8.014	-15.617	1.00	8.01	SEG1
ATOM	99	NE2	HIS	8	-14.828	8.952	-14.213	1.00	7.43	SEG1
ATOM	100	C	HIS	8	-18.520	11.331	-11.664	1.00	4.62	SEG1
ATOM	101	O	HIS	8	-18.996	12.104	-10.854	1.00	4.56	SEG1
ATOM	102	N	GLU	9	-18.163	10.118	-11.327	1.00	4.40	SEG1
ATOM	103	HN	GLU	9	-17.783	9.518	-12.002	1.00	4.79	SEG1
ATOM	104	CA	GLU	9	-18.332	9.637	-9.922	1.00	3.95	SEG1
ATOM	105	HA	GLU	9	-19.165	10.132	-9.449	1.00	4.17	SEG1
ATOM	106	CB	GLU	9	-18.618	8.140	-10.058	1.00	4.47	SEG1
ATOM	107	HB1	GLU	9	-18.580	7.675	-9.084	1.00	4.63	SEG1
ATOM	108	HB2	GLU	9	-17.877	7.689	-10.703	1.00	4.60	SEG1
ATOM	109	CG	GLU	9	-20.011	7.939	-10.661	1.00	5.14	SEG1
ATOM	110	HG1	GLU	9	-20.102	6.931	-11.038	1.00	5.23	SEG1
ATOM	111	HG2	GLU	9	-20.157	8.642	-11.469	1.00	5.30	SEG1
ATOM	112	CD	GLU	9	-21.069	8.175	-9.582	1.00	5.96	SEG1
ATOM	113	OE1	GLU	9	-22.112	8.718	-9.908	1.00	6.30	SEG1
ATOM	114	OE2	GLU	9	-20.819	7.806	-8.446	1.00	6.51	SEG1
ATOM	115	C	GLU	9	-17.043	9.872	-9.127	1.00	3.32	SEG1
ATOM	116	O	GLU	9	-15.955	9.639	-9.621	1.00	3.53	SEG1
ATOM	117	N	GLU	10	-17.155	10.329	-7.902	1.00	2.92	SEG1
ATOM	118	HN	GLU	10	-18.042	10.508	-7.525	1.00	3.15	SEG1
ATOM	119	CA	GLU	10	-15.930	10.572	-7.084	1.00	2.63	SEG1
ATOM	120	HA	GLU	10	-15.122	10.912	-7.713	1.00	2.97	SEG1
ATOM	121	CB	GLU	10	-16.306	11.667	-6.084	1.00	3.13	SEG1
ATOM	122	HB1	GLU	10	-15.459	11.879	-5.449	1.00	3.26	SEG1
ATOM	123	HB2	GLU	10	-17.134	11.333	-5.477	1.00	3.47	SEG1
ATOM	124	CG	GLU	10	-16.703	12.940	-6.840	1.00	3.83	SEG1
ATOM	125	HG1	GLU	10	-17.631	12.768	-7.365	1.00	4.16	SEG1
ATOM	126	HG2	GLU	10	-15.930	13.189	-7.552	1.00	4.02	SEG1
ATOM	127	CD	GLU	10	-16.888	14.107	-5.858	1.00	4.51	SEG1
ATOM	128	OE1	GLU	10	-16.595	13.936	-4.682	1.00	5.01	SEG1
ATOM	129	OE2	GLU	10	-17.319	15.159	-6.302	1.00	4.85	SEG1
ATOM	130	C	GLU	10	-15.531	9.290	-6.355	1.00	2.02	SEG1
ATOM	131	O	GLU	10	-16.296	8.732	-5.591	1.00	2.17	SEG1
ATOM	132	N	TRP	11	-14.339	8.818	-6.599	1.00	1.56	SEG1
ATOM	133	HN	TRP	11	-13.749	9.288	-7.225	1.00	1.73	SEG1
ATOM	134	CA	TRP	11	-13.868	7.567	-5.940	1.00	1.15	SEG1
ATOM	135	HA	TRP	11	-14.611	7.214	-5.241	1.00	1.26	SEG1
ATOM	136	CB	TRP	11	-13.737	6.567	-7.116	1.00	1.36	SEG1
ATOM	137	HB1	TRP	11	-13.492	7.113	-8.017	1.00	1.88	SEG1
ATOM	138	HB2	TRP	11	-14.682	6.065	-7.257	1.00	1.58	SEG1
ATOM	139	CG	TRP	11	-12.677	5.536	-6.862	1.00	1.30	SEG1
ATOM	140	CD1	TRP	11	-11.365	5.717	-7.112	1.00	2.28	SEG1
ATOM	141	HD1	TRP	11	-10.924	6.614	-7.517	1.00	3.09	SEG1

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ATOM	142	COO	TRP	11	-12.809	4.189	-6.326	1.00	0.69	SEG1
ATOM	143	NE1	TRP	11	-10.683	4.571	-6.754	1.00	2.24	SEG1
ATOM	144	HE1	TPP	11	-9.724	4.445	-6.846	1.00	2.95	SEG1
ATOM	145	CE2	TRP	11	-11.525	3.600	-6.174	1.00	1.19	SEG1
ATOM	146	CE3	TRP	11	-13.901	3.430	-5.884	1.00	1.24	SEG1
ATOM	147	HE3	TPP	11	-14.889	3.854	-5.920	1.00	1.82	SEG1
ATOM	148	CE2	TPP	11	-11.331	2.304	-5.601	1.00	1.05	SEG1
ATOM	149	HE2	TPP	11	-10.339	1.878	-5.771	1.00	1.53	SEG1
ATOM	150	CE3	TPP	11	-13.712	2.125	-5.406	1.00	1.72	SEG1
ATOM	151	HE3	TPP	11	-14.560	1.550	-5.068	1.00	2.64	SEG1
ATOM	152	CH2	TRP	11	-12.428	1.563	-5.365	1.00	1.25	SEG1
ATOM	153	HH2	TRP	11	-12.289	0.559	-4.995	1.00	1.70	SEG1
ATOM	154	C	TPP	11	-12.535	7.832	-5.112	1.00	1.07	SEG1
ATOM	155	O	TPP	11	-11.569	8.269	-5.807	1.00	1.43	SEG1
ATOM	156	N	VAL	12	-12.488	7.557	-3.933	1.00	0.90	SEG1
ATOM	157	HN	VAL	12	-13.283	7.199	-3.483	1.00	1.07	SEG1
ATOM	158	CA	VAL	12	-11.230	7.774	-3.153	1.00	0.81	SEG1
ATOM	159	HA	VAL	12	-10.374	7.754	-3.809	1.00	0.90	SEG1
ATOM	160	CB	VAL	12	-11.366	9.163	-2.513	1.00	1.11	SEG1
ATOM	161	HB	VAL	12	-10.500	9.347	-1.893	1.00	1.25	SEG1
ATOM	162	CG1	VAL	12	-11.445	10.238	-3.600	1.00	1.39	SEG1
ATOM	163	HG11	VAL	12	-11.487	11.113	-3.136	1.00	1.88	SEG1
ATOM	164	HG12	VAL	12	-12.332	10.087	-4.195	1.00	1.88	SEG1
ATOM	165	HG13	VAL	12	-10.571	10.178	-4.231	1.00	1.65	SEG1
ATOM	166	CG2	VAL	12	-12.632	9.223	-1.651	1.00	1.61	SEG1
ATOM	167	HG21	VAL	12	-12.669	10.169	-1.132	1.00	2.15	SEG1
ATOM	168	HG22	VAL	12	-12.616	8.419	-0.931	1.00	1.97	SEG1
ATOM	169	HG23	VAL	12	-13.504	9.125	-2.181	1.00	1.99	SEG1
ATOM	170	C	VAL	12	-11.095	6.703	-2.063	1.00	0.82	SEG1
ATOM	171	O	VAL	12	-10.758	7.001	-0.932	1.00	1.69	SEG1
ATOM	172	N	GLY	13	-11.375	5.466	-2.388	1.00	0.62	SEG1
ATOM	173	HN	GLY	13	-11.660	5.252	-3.302	1.00	1.15	SEG1
ATOM	174	CA	GLY	13	-11.287	4.378	-1.367	1.00	0.89	SEG1
ATOM	175	HA1	GLY	13	-11.623	3.451	-1.805	1.00	1.13	SEG1
ATOM	176	HA2	GLY	13	-11.922	4.626	-0.529	1.00	1.25	SEG1
ATOM	177	C	GLY	13	-9.848	4.206	-0.874	1.00	0.68	SEG1
ATOM	178	O	GLY	13	-8.900	4.566	-1.544	1.00	0.70	SEG1
ATOM	179	N	SER	14	-9.692	3.645	0.299	1.00	0.57	SEG1
ATOM	180	HN	SER	14	-10.481	3.363	0.809	1.00	0.64	SEG1
ATOM	181	CA	SER	14	-8.330	3.421	0.867	1.00	0.46	SEG1
ATOM	182	HA	SER	14	-7.570	3.656	0.139	1.00	0.47	SEG1
ATOM	183	CB	SER	14	-8.235	4.373	2.057	1.00	0.63	SEG1
ATOM	184	HB1	SER	14	-8.473	5.378	1.733	1.00	1.22	SEG1
ATOM	185	HB2	SER	14	-7.235	4.356	2.457	1.00	1.22	SEG1
ATOM	186	DG	SER	14	-9.150	3.958	3.063	1.00	1.27	SEG1
ATOM	187	HG	SER	14	-8.910	4.396	3.882	1.00	1.63	SEG1
ATOM	188	C	SER	14	-8.206	1.967	1.331	1.00	0.35	SEG1
ATOM	189	O	SER	14	-9.198	1.309	1.587	1.00	0.36	SEG1
ATOM	190	N	ALA	15	-7.004	1.456	1.433	1.00	0.31	SEG1
ATOM	191	HN	ALA	15	-6.220	2.001	1.215	1.00	0.37	SEG1
ATOM	192	CA	ALA	15	-6.830	0.036	1.869	1.00	0.26	SEG1
ATOM	193	HA	ALA	15	-7.719	-0.320	2.363	1.00	0.26	SEG1
ATOM	194	CB	ALA	15	-6.611	-0.741	0.573	1.00	0.33	SEG1
ATOM	195	HB1	ALA	15	-7.409	-0.518	-0.120	1.00	1.05	SEG1
ATOM	196	HB2	ALA	15	-6.603	-1.800	0.785	1.00	1.09	SEG1
ATOM	197	HB3	ALA	15	-5.664	-0.454	0.138	1.00	1.01	SEG1
ATOM	198	C	ALA	15	-5.612	-0.112	2.785	1.00	0.24	SEG1
ATOM	199	O	ALA	15	-4.700	0.693	2.759	1.00	0.25	SEG1
ATOM	200	N	TYR	16	-5.590	-1.151	3.584	1.00	0.24	SEG1
ATOM	201	HN	TYR	16	-6.336	-1.788	3.574	1.00	0.25	SEG1
ATOM	202	CA	TYR	16	-4.431	-1.381	4.498	1.00	0.24	SEG1
ATOM	203	HA	TYR	16	-3.769	-0.529	4.495	1.00	0.25	SEG1
ATOM	204	CB	TYR	16	-5.045	-1.561	5.886	1.00	0.28	SEG1
ATOM	205	HB1	TYR	16	-4.302	-1.960	6.560	1.00	0.30	SEG1
ATOM	206	HB2	TYR	16	-5.880	-2.244	5.825	1.00	0.29	SEG1
ATOM	207	CG	TYR	16	-5.524	-0.224	6.401	1.00	0.30	SEG1
ATOM	208	CD1	TYR	16	-6.885	0.100	6.365	1.00	1.20	SEG1
ATOM	209	HD1	TYR	16	-7.597	-0.607	5.967	1.00	2.11	SEG1
ATOM	210	CD2	TYR	16	-4.601	0.693	6.916	1.00	1.30	SEG1
ATOM	211	HD2	TYR	16	-3.550	0.442	6.943	1.00	2.22	SEG1
ATOM	212	CE1	TYR	16	-7.322	1.340	6.845	1.00	1.20	SEG1
ATOM	213	HE1	TYR	16	-8.372	1.591	6.818	1.00	2.12	SEG1
ATOM	214	CE2	TYR	16	-5.036	1.933	7.395	1.00	1.32	SEG1
ATOM	215	HE2	TYR	16	-4.323	2.639	7.792	1.00	2.24	SEG1
ATOM	216	CZ	TYR	16	-6.397	2.259	7.361	1.00	0.37	SEG1
ATOM	217	OH	TYR	16	-6.826	3.483	7.839	1.00	0.42	SEG1
ATOM	218	HH	TYR	16	-7.602	3.747	7.339	1.00	1.02	SEG1

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FIG. 2 (4 of 35)

ATOM	219	C	TYR	16	-3.678	-2.645	4.068	1.00	0.23	SEG1
ATOM	220	O	TYR	16	-4.273	-3.675	3.812	1.00	0.25	SEG1
ATOM	221	N	LEU	17	-2.377	-2.565	3.978	1.00	0.23	SEG1
ATOM	222	HN	LEU	17	-1.930	-1.721	4.183	1.00	0.23	SEG1
ATOM	223	CA	LEU	17	-1.570	-3.747	3.551	1.00	0.23	SEG1
ATOM	224	HA	LEU	17	-2.217	-4.569	3.299	1.00	0.24	SEG1
ATOM	225	CB	LEU	17	-0.815	-3.278	2.356	1.00	0.25	SEG1
ATOM	226	HB1	LEU	17	0.169	-3.723	2.294	1.00	0.26	SEG1
ATOM	227	HB2	LEU	17	-0.724	-2.102	2.326	1.00	0.25	SEG1
ATOM	228	CG	LEU	17	-1.577	-3.703	1.051	1.00	0.25	SEG1
ATOM	229	HG	LEU	17	-1.713	-4.777	1.063	1.00	0.26	SEG1
ATOM	230	CD1	LEU	17	-2.943	-3.009	1.014	1.00	0.25	SEG1
ATOM	231	HD11	LEU	17	-3.272	-2.919	-0.010	1.00	1.04	SEG1
ATOM	232	HD12	LEU	17	-2.859	-2.026	1.453	1.00	1.04	SEG1
ATOM	233	HD13	LEU	17	-3.659	-3.593	1.573	1.00	1.03	SEG1
ATOM	234	CD2	LEU	17	-0.775	-3.299	-0.188	1.00	0.28	SEG1
ATOM	235	HD21	LEU	17	-1.452	-3.080	-0.999	1.00	1.03	SEG1
ATOM	236	HD22	LEU	17	-0.120	-4.109	-0.473	1.00	1.07	SEG1
ATOM	237	HD23	LEU	17	-0.187	-2.421	0.037	1.00	1.01	SEG1
ATOM	238	C	LEU	17	-0.587	-4.159	4.646	1.00	0.24	SEG1
ATOM	239	O	LEU	17	-0.008	-3.327	5.318	1.00	0.24	SEG1
ATOM	240	N	PHE	18	-0.397	-5.443	4.826	1.00	0.26	SEG1
ATOM	241	HN	PHE	18	-0.877	-6.090	4.269	1.00	0.28	SEG1
ATOM	242	CA	PHE	18	0.549	-5.923	5.878	1.00	0.28	SEG1
ATOM	243	HA	PHE	18	0.825	-5.116	6.536	1.00	0.26	SEG1
ATOM	244	CB	PHE	18	-0.221	-6.998	6.657	1.00	0.30	SEG1
ATOM	245	HB1	PHE	18	0.392	-7.355	7.471	1.00	0.31	SEG1
ATOM	246	HB2	PHE	18	-0.447	-7.820	5.996	1.00	0.31	SEG1
ATOM	247	CG	PHE	18	-1.514	-6.433	7.214	1.00	0.29	SEG1
ATOM	248	CD1	PHE	18	-1.550	-5.139	7.750	1.00	1.21	SEG1
ATOM	249	HD1	PHE	18	-0.652	-4.540	7.764	1.00	2.13	SEG1
ATOM	250	CD2	PHE	18	-2.678	-7.210	7.191	1.00	1.27	SEG1
ATOM	251	HD2	PHE	18	-2.652	-8.208	6.779	1.00	2.18	SEG1
ATOM	252	CE1	PHE	18	-2.745	-4.624	8.261	1.00	1.20	SEG1
ATOM	253	HE1	PHE	18	-2.770	-3.626	8.674	1.00	2.11	SEG1
ATOM	254	CE2	PHE	18	-3.874	-6.695	7.703	1.00	1.29	SEG1
ATOM	255	HE2	PHE	18	-4.772	-7.296	7.685	1.00	2.22	SEG1
ATOM	256	CZ	PHE	18	-3.907	-5.402	8.238	1.00	0.32	SEG1
ATOM	257	HZ	PHE	18	-4.828	-5.004	8.631	1.00	0.35	SEG1
ATOM	258	C	PHE	18	1.787	-6.535	5.226	1.00	0.30	SEG1
ATOM	259	O	PHE	18	1.681	-7.379	4.360	1.00	0.33	SEG1
ATOM	260	N	VAL	19	2.956	-6.123	5.641	1.00	0.29	SEG1
ATOM	261	HN	VAL	19	3.013	-5.443	6.346	1.00	0.27	SEG1
ATOM	262	CA	VAL	19	4.206	-6.689	5.048	1.00	0.32	SEG1
ATOM	263	HA	VAL	19	3.970	-7.353	4.230	1.00	0.35	SEG1
ATOM	264	CB	VAL	19	4.993	-5.478	4.535	1.00	0.33	SEG1
ATOM	265	HB	VAL	19	5.223	-4.826	5.367	1.00	0.31	SEG1
ATOM	266	CG1	VAL	19	6.300	-5.953	3.891	1.00	0.37	SEG1
ATOM	267	HG11	VAL	19	6.080	-6.458	2.962	1.00	1.07	SEG1
ATOM	268	HG12	VAL	19	6.804	-6.635	4.560	1.00	1.07	SEG1
ATOM	269	HG13	VAL	19	6.936	-5.102	3.696	1.00	1.08	SEG1
ATOM	270	CG2	VAL	19	4.157	-4.712	3.500	1.00	0.36	SEG1
ATOM	271	HG21	VAL	19	4.520	-4.930	2.507	1.00	1.03	SEG1
ATOM	272	HG22	VAL	19	4.240	-3.652	3.687	1.00	1.07	SEG1
ATOM	273	HG23	VAL	19	3.121	-5.010	3.578	1.00	1.13	SEG1
ATOM	274	C	VAL	19	5.001	-7.426	6.128	1.00	0.32	SEG1
ATOM	275	O	VAL	19	5.376	-6.844	7.131	1.00	0.31	SEG1
ATOM	276	N	GLU	20	5.260	-8.698	5.937	1.00	0.36	SEG1
ATOM	277	HN	GLU	20	4.945	-9.149	5.122	1.00	0.38	SEG1
ATOM	278	CA	GLU	20	6.035	-9.459	6.969	1.00	0.38	SEG1
ATOM	279	HA	GLU	20	6.640	-8.784	7.553	1.00	0.38	SEG1
ATOM	280	CB	GLU	20	4.968	-10.101	7.858	1.00	0.41	SEG1
ATOM	281	HB1	GLU	20	4.576	-10.981	7.372	1.00	0.96	SEG1
ATOM	282	HB2	GLU	20	4.167	-9.393	8.025	1.00	0.89	SEG1
ATOM	283	CG	GLU	20	5.581	-10.495	9.206	1.00	1.26	SEG1
ATOM	284	HG1	GLU	20	5.966	-9.612	9.693	1.00	1.79	SEG1
ATOM	285	HG2	GLU	20	6.387	-11.193	9.045	1.00	1.83	SEG1
ATOM	286	CD	GLU	20	4.515	-11.141	10.106	1.00	1.26	SEG1
ATOM	287	OE1	GLU	20	4.890	-11.676	11.136	1.00	1.44	SEG1
ATOM	288	OE2	GLU	20	3.343	-11.084	9.758	1.00	1.82	SEG1
ATOM	289	C	GLU	20	6.920	-10.527	6.305	1.00	0.41	SEG1
ATOM	290	O	GLU	20	6.590	-11.059	5.264	1.00	0.48	SEG1
ATOM	291	N	SER	21	8.045	-10.834	6.906	1.00	0.42	SEG1
ATOM	292	HN	SER	21	8.286	-10.383	7.743	1.00	0.44	SEG1
ATOM	293	CA	SER	21	8.971	-11.860	6.324	1.00	0.48	SEG1
ATOM	294	HA	SER	21	8.796	-11.968	5.265	1.00	0.50	SEG1
ATOM	295	CB	SER	21	10.372	-11.292	5.554	1.00	0.52	SEG1

ATOM	296	HB1	SER	21	10.474	-10.360	6.014	1.00	0.55	SEG1
ATOM	297	HB2	SER	21	11.109	-11.992	6.197	1.00	0.54	SEG1
ATOM	298	OG	SER	21	10.572	-11.073	7.945	1.00	0.62	SEG1
ATOM	299	4G	SER	21	10.391	-11.893	8.405	1.00	1.01	SEG1
ATOM	300	C	SER	21	8.820	-13.120	7.030	1.00	0.56	SEG1
ATOM	301	O	SER	21	9.613	-14.118	6.856	1.00	1.22	SEG1
ATOM	302	N	SER	22	7.817	-13.178	7.663	1.00	0.86	SEG1
ATOM	303	HN	SER	22	7.193	-12.643	8.016	1.00	1.44	SEG1
ATOM	304	CA	SER	22	7.615	-14.675	8.592	1.00	0.95	SEG1
ATOM	305	HA	SER	22	6.749	-14.607	9.232	1.00	1.07	SEG1
ATOM	306	CB	SER	22	7.374	-15.735	7.512	1.00	1.04	SEG1
ATOM	307	HB1	SER	22	8.297	-16.265	7.318	1.00	1.13	SEG1
ATOM	308	HB2	SER	22	7.039	-15.260	6.605	1.00	1.30	SEG1
ATOM	309	OG	SER	22	6.376	-16.643	7.960	1.00	1.42	SEG1
ATOM	310	HG	SER	22	6.500	-17.474	7.495	1.00	1.68	SEG1
ATOM	311	C	SER	22	8.855	-15.024	9.424	1.00	0.91	SEG1
ATOM	312	O	SER	22	9.120	-16.180	9.698	1.00	0.94	SEG1
ATOM	313	N	LEU	23	9.604	-14.033	9.841	1.00	0.96	SEG1
ATOM	314	HN	LEU	23	9.362	-13.111	9.617	1.00	1.04	SEG1
ATOM	315	CA	LEU	23	10.819	-14.303	10.673	1.00	1.03	SEG1
ATOM	316	HA	LEU	23	10.973	-15.365	10.782	1.00	1.20	SEG1
ATOM	317	CB	LEU	23	11.986	-13.678	9.904	1.00	0.94	SEG1
ATOM	318	HB1	LEU	23	12.861	-13.658	10.537	1.00	1.14	SEG1
ATOM	319	HB2	LEU	23	11.729	-12.670	9.616	1.00	1.11	SEG1
ATOM	320	CG	LEU	23	12.286	-14.508	8.654	1.00	1.17	SEG1
ATOM	321	HG	LEU	23	11.371	-14.691	8.112	1.00	1.54	SEG1
ATOM	322	CD1	LEU	23	13.270	-13.748	7.762	1.00	1.51	SEG1
ATOM	323	HD11	LEU	23	12.805	-12.843	7.401	1.00	1.96	SEG1
ATOM	324	HD12	LEU	23	13.547	-14.368	6.921	1.00	2.02	SEG1
ATOM	325	HD13	LEU	23	14.153	-13.497	8.331	1.00	1.83	SEG1
ATOM	326	CD2	LEU	23	12.914	-15.839	9.073	1.00	1.52	SEG1
ATOM	327	HD21	LEU	23	13.458	-16.258	8.240	1.00	1.93	SEG1
ATOM	328	HD22	LEU	23	12.138	-16.524	9.378	1.00	2.01	SEG1
ATOM	329	HD23	LEU	23	13.592	-15.672	9.898	1.00	1.88	SEG1
ATOM	330	C	LEU	23	10.668	-13.638	12.043	1.00	1.14	SEG1
ATOM	331	O	LEU	23	10.726	-12.430	12.165	1.00	1.44	SEG1
ATOM	332	N	ASP	24	10.479	-14.422	13.073	1.00	1.17	SEG1
ATOM	333	HN	ASP	24	10.438	-15.393	12.943	1.00	1.26	SEG1
ATOM	334	CA	ASP	24	10.324	-13.844	14.444	1.00	1.33	SEG1
ATOM	335	HA	ASP	24	9.561	-13.082	14.444	1.00	1.51	SEG1
ATOM	336	CB	ASP	24	9.887	-15.016	15.327	1.00	1.68	SEG1
ATOM	337	HB1	ASP	24	9.925	-14.718	16.365	1.00	1.96	SEG1
ATOM	338	HB2	ASP	24	10.552	-15.851	15.169	1.00	1.96	SEG1
ATOM	339	CG	ASP	24	8.455	-15.430	14.972	1.00	1.87	SEG1
ATOM	340	OD1	ASP	24	8.137	-16.594	15.151	1.00	2.18	SEG1
ATOM	341	OD2	ASP	24	7.700	-14.577	14.532	1.00	2.42	SEG1
ATOM	342	C	ASP	24	11.654	-13.263	14.943	1.00	1.21	SEG1
ATOM	343	O	ASP	24	11.680	-12.453	15.850	1.00	1.73	SEG1
ATOM	344	N	LYS	25	12.757	-13.677	14.367	1.00	1.40	SEG1
ATOM	345	HN	LYS	25	12.714	-14.337	13.644	1.00	1.84	SEG1
ATOM	346	CA	LYS	25	14.086	-13.157	14.816	1.00	1.71	SEG1
ATOM	347	HA	LYS	25	14.109	-13.063	15.889	1.00	1.98	SEG1
ATOM	348	CB	LYS	25	15.096	-14.216	14.368	1.00	2.08	SEG1
ATOM	349	HB1	LYS	25	16.098	-13.842	14.519	1.00	2.36	SEG1
ATOM	350	HB2	LYS	25	14.947	-14.430	13.320	1.00	1.98	SEG1
ATOM	351	CG	LYS	25	14.905	-15.498	15.181	1.00	2.37	SEG1
ATOM	352	HG1	LYS	25	13.906	-15.877	15.028	1.00	2.26	SEG1
ATOM	353	HG2	LYS	25	15.054	-15.286	16.230	1.00	2.63	SEG1
ATOM	354	CD	LYS	25	15.922	-16.545	14.720	1.00	2.85	SEG1
ATOM	355	HD1	LYS	25	16.921	-16.166	14.874	1.00	3.30	SEG1
ATOM	356	HD2	LYS	25	15.774	-16.753	13.670	1.00	2.96	SEG1
ATOM	357	CE	LYS	25	15.738	-17.833	15.527	1.00	3.10	SEG1
ATOM	358	HE1	LYS	25	14.780	-18.279	15.310	1.00	3.12	SEG1
ATOM	359	HE2	LYS	25	15.831	-17.629	16.584	1.00	3.39	SEG1
ATOM	360	NZ	LYS	25	16.834	-18.732	15.069	1.00	3.59	SEG1
ATOM	361	H21	LYS	25	17.748	-18.362	15.398	1.00	3.96	SEG1
ATOM	362	H22	LYS	25	16.833	-18.778	14.029	1.00	3.70	SEG1
ATOM	363	H23	LYS	25	16.688	-19.684	15.458	1.00	3.91	SEG1
ATOM	364	C	LYS	25	14.414	-11.811	14.152	1.00	1.50	SEG1
ATOM	365	O	LYS	25	15.402	-11.183	14.489	1.00	1.60	SEG1
ATOM	366	N	VAL	26	13.614	-11.362	13.211	1.00	1.35	SEG1
ATOM	367	HN	VAL	26	12.826	-11.876	12.943	1.00	1.43	SEG1
ATOM	368	CA	VAL	26	13.912	-10.062	12.538	1.00	1.21	SEG1
ATOM	369	HA	VAL	26	14.886	-9.702	12.832	1.00	1.29	SEG1
ATOM	370	CB	VAL	26	13.903	-10.380	11.037	1.00	1.42	SEG1
ATOM	371	HB	VAL	26	12.932	-10.761	10.756	1.00	1.62	SEG1
ATOM	372	CG1	VAL	26	14.200	-9.108	10.237	1.00	1.89	SEG1

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ATOM	373	HG11	VAL	26	13.971	-9.278	9.195	1.00	2.27	SEG1
ATOM	374	HG12	VAL	26	15.244	-8.954	10.340	1.00	2.38	SEG1
ATOM	375	HG13	VAL	26	13.594	-8.297	10.611	1.00	2.31	SEG1
ATOM	376	CG2	VAL	26	14.973	-11.431	10.728	1.00	2.08	SEG1
ATOM	377	HG21	VAL	26	15.828	-11.275	11.370	1.00	2.45	SEG1
ATOM	378	HG22	VAL	26	15.278	-11.342	9.696	1.00	2.51	SEG1
ATOM	379	HG23	VAL	26	14.569	-12.418	10.900	1.00	2.57	SEG1
ATOM	380	C	VAL	26	12.836	-9.026	12.877	1.00	1.02	SEG1
ATOM	381	O	VAL	26	11.670	-9.210	12.577	1.00	1.13	SEG1
ATOM	382	N	VAL	27	13.225	-7.935	13.484	1.00	0.99	SEG1
ATOM	383	HN	VAL	27	14.171	-7.811	13.703	1.00	1.14	SEG1
ATOM	384	CA	VAL	27	12.232	-6.873	13.827	1.00	1.00	SEG1
ATOM	385	HA	VAL	27	11.271	-7.317	14.042	1.00	1.11	SEG1
ATOM	386	CB	VAL	27	12.783	-6.182	15.085	1.00	1.28	SEG1
ATOM	387	HB	VAL	27	12.091	-5.413	15.399	1.00	1.71	SEG1
ATOM	388	CG1	VAL	27	12.928	-7.213	16.209	1.00	1.74	SEG1
ATOM	389	HG11	VAL	27	11.949	-7.521	16.544	1.00	2.16	SEG1
ATOM	390	HG12	VAL	27	13.470	-6.773	17.033	1.00	2.26	SEG1
ATOM	391	HG13	VAL	27	13.469	-8.072	15.841	1.00	2.14	SEG1
ATOM	392	CG2	VAL	27	14.154	-5.547	14.792	1.00	1.90	SEG1
ATOM	393	HG21	VAL	27	14.856	-5.818	15.568	1.00	2.40	SEG1
ATOM	394	HG22	VAL	27	14.051	-4.473	14.763	1.00	2.42	SEG1
ATOM	395	HG23	VAL	27	14.521	-5.896	13.838	1.00	2.26	SEG1
ATOM	396	C	VAL	27	12.112	-5.891	12.655	1.00	0.83	SEG1
ATOM	397	O	VAL	27	12.578	-4.769	12.717	1.00	0.80	SEG1
ATOM	398	N	LEU	28	11.497	-6.319	11.581	1.00	0.77	SEG1
ATOM	399	HN	LEU	28	11.142	-7.236	11.559	1.00	0.85	SEG1
ATOM	400	CA	LEU	28	11.353	-5.429	10.385	1.00	0.65	SEG1
ATOM	401	HA	LEU	28	12.323	-5.176	9.987	1.00	0.73	SEG1
ATOM	402	CB	LEU	28	10.571	-6.248	9.359	1.00	0.66	SEG1
ATOM	403	HB1	LEU	28	10.277	-5.610	8.539	1.00	0.81	SEG1
ATOM	404	HB2	LEU	28	9.689	-6.660	9.829	1.00	0.72	SEG1
ATOM	405	CG	LEU	28	11.445	-7.384	8.832	1.00	1.04	SEG1
ATOM	406	HG	LEU	28	11.877	-7.921	9.664	1.00	1.45	SEG1
ATOM	407	CD1	LEU	28	10.588	-8.336	8.001	1.00	1.13	SEG1
ATOM	408	HD11	LEU	28	9.882	-7.766	7.415	1.00	1.40	SEG1
ATOM	409	HD12	LEU	28	10.052	-9.005	8.658	1.00	1.73	SEG1
ATOM	410	HD13	LEU	28	11.222	-8.910	7.342	1.00	1.54	SEG1
ATOM	411	CD2	LEU	28	12.560	-6.805	7.959	1.00	1.38	SEG1
ATOM	412	HD21	LEU	28	12.188	-5.943	7.425	1.00	1.93	SEG1
ATOM	413	HD22	LEU	28	12.889	-7.553	7.253	1.00	1.66	SEG1
ATOM	414	HD23	LEU	28	13.391	-6.510	8.584	1.00	1.78	SEG1
ATOM	415	C	LEU	28	10.576	-4.160	10.737	1.00	0.52	SEG1
ATOM	416	O	LEU	28	10.902	-3.082	10.279	1.00	0.49	SEG1
ATOM	417	N	SER	29	9.547	-4.279	11.537	1.00	0.53	SEG1
ATOM	418	HN	SER	29	9.302	-5.163	11.885	1.00	0.61	SEG1
ATOM	419	CA	SER	29	8.735	-3.075	11.909	1.00	0.50	SEG1
ATOM	420	HA	SER	29	8.245	-2.669	11.037	1.00	0.45	SEG1
ATOM	421	CB	SER	29	7.690	-3.587	12.903	1.00	0.64	SEG1
ATOM	422	HB1	SER	29	7.195	-4.456	12.489	1.00	0.68	SEG1
ATOM	423	HB2	SER	29	6.961	-2.817	13.090	1.00	0.69	SEG1
ATOM	424	OG	SER	29	8.332	-3.930	14.125	1.00	0.74	SEG1
ATOM	425	HG	SER	29	7.660	-4.234	14.739	1.00	1.18	SEG1
ATOM	426	C	SER	29	9.622	-2.014	12.569	1.00	0.51	SEG1
ATOM	427	O	SER	29	9.388	-0.827	12.436	1.00	0.47	SEG1
ATOM	428	N	ASP	30	10.640	-2.436	13.274	1.00	0.61	SEG1
ATOM	429	HN	ASP	30	10.805	-3.399	13.361	1.00	0.66	SEG1
ATOM	430	CA	ASP	30	11.555	-1.459	13.946	1.00	0.70	SEG1
ATOM	431	HA	ASP	30	11.012	-0.874	14.672	1.00	0.74	SEG1
ATOM	432	CB	ASP	30	12.614	-2.315	14.651	1.00	0.86	SEG1
ATOM	433	HB1	ASP	30	13.205	-2.833	13.911	1.00	1.25	SEG1
ATOM	434	HB2	ASP	30	12.123	-3.037	15.288	1.00	0.89	SEG1
ATOM	435	CG	ASP	30	13.531	-1.426	15.500	1.00	1.76	SEG1
ATOM	436	OD1	ASP	30	13.699	-0.267	15.156	1.00	2.44	SEG1
ATOM	437	OD2	ASP	30	14.049	-1.922	16.488	1.00	2.38	SEG1
ATOM	438	C	ASP	30	12.203	-0.543	12.901	1.00	0.66	SEG1
ATOM	439	O	ASP	30	12.366	0.643	13.120	1.00	0.69	SEG1
ATOM	440	N	ALA	31	12.580	-1.089	11.774	1.00	0.65	SEG1
ATOM	441	HN	ALA	31	12.442	-2.049	11.629	1.00	0.64	SEG1
ATOM	442	CA	ALA	31	13.229	-0.259	10.710	1.00	0.69	SEG1
ATOM	443	HA	ALA	31	14.158	0.154	11.072	1.00	0.81	SEG1
ATOM	444	CB	ALA	31	13.506	-1.226	9.556	1.00	0.73	SEG1
ATOM	445	HB1	ALA	31	13.622	-2.227	9.945	1.00	1.33	SEG1
ATOM	446	HB2	ALA	31	14.413	-0.930	9.048	1.00	1.15	SEG1
ATOM	447	HB3	ALA	31	12.681	-1.203	8.860	1.00	1.27	SEG1
ATOM	448	C	ALA	31	12.290	0.862	10.254	1.00	0.59	SEG1
ATOM	449	O	ALA	31	12.716	1.974	10.014	1.00	0.66	SEG1

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ATCM	527	C	GLN	36	13.784	7.724	7.081	1.00	0.44	SEG1
ATCM	528	O	GLN	36	13.699	7.773	5.868	1.00	0.45	SEG1
ATCM	529	N	GLN	37	13.641	6.602	7.734	1.00	0.41	SEG1
ATCM	530	HN	GLN	37	13.704	6.599	8.714	1.00	0.41	SEG1
ATCM	531	CA	GLN	37	13.363	5.331	7.003	1.00	0.40	SEG1
ATCM	532	HA	GLN	37	14.080	5.189	6.213	1.00	0.45	SEG1
ATCM	533	CB	GLN	37	13.518	4.228	9.052	1.00	0.41	SEG1
ATCM	534	HB1	GLN	37	13.168	3.290	7.649	1.00	0.42	SEG1
ATCM	535	HB2	GLN	37	12.944	4.483	8.932	1.00	0.38	SEG1
ATCM	536	CG	GLN	37	14.999	4.103	8.424	1.00	0.48	SEG1
ATCM	537	HG1	GLN	37	15.356	5.048	8.803	1.00	0.60	SEG1
ATCM	538	HG2	GLN	37	15.566	3.830	7.546	1.00	0.64	SEG1
ATCM	539	CD	GLN	37	15.178	3.027	9.497	1.00	0.50	SEG1
ATCM	540	OE1	GLN	37	14.775	3.208	10.626	1.00	1.10	SEG1
ATCM	541	NE2	GLN	37	15.773	1.907	9.193	1.00	0.90	SEG1
ATCM	542	HE21	GLN	37	16.103	1.758	8.282	1.00	1.44	SEG1
ATCM	543	HE22	GLN	37	15.889	1.213	9.876	1.00	0.99	SEG1
ATCM	544	C	GLN	37	11.950	5.366	6.422	1.00	0.36	SEG1
ATCM	545	O	GLN	37	11.706	4.870	5.340	1.00	0.37	SEG1
ATCM	546	N	LYS	38	11.022	5.968	7.125	1.00	0.33	SEG1
ATCM	547	HN	LYS	38	11.253	6.372	7.987	1.00	0.33	SEG1
ATCM	548	CA	LYS	38	9.620	6.055	6.604	1.00	0.31	SEG1
ATCM	549	HA	LYS	38	9.196	5.071	6.484	1.00	0.32	SEG1
ATCM	550	CB	LYS	38	8.849	6.839	7.672	1.00	0.33	SEG1
ATCM	551	HB1	LYS	38	9.294	7.815	7.792	1.00	0.35	SEG1
ATCM	552	HB2	LYS	38	8.894	6.306	8.610	1.00	0.34	SEG1
ATCM	553	CG	LYS	38	7.388	6.999	7.251	1.00	0.36	SEG1
ATCM	554	HG1	LYS	38	6.937	6.024	7.138	1.00	0.37	SEG1
ATCM	555	HG2	LYS	38	7.338	7.532	6.312	1.00	0.36	SEG1
ATCM	556	CD	LYS	38	6.635	7.784	8.328	1.00	0.46	SEG1
ATCM	557	HD1	LYS	38	7.088	8.756	8.450	1.00	1.19	SEG1
ATCM	558	HD2	LYS	38	6.682	7.244	9.264	1.00	1.04	SEG1
ATCM	559	CE	LYS	38	5.175	7.953	7.910	1.00	1.07	SEG1
ATCM	560	HE1	LYS	38	4.696	6.991	7.826	1.00	1.63	SEG1
ATCM	561	HE2	LYS	38	5.113	8.490	6.973	1.00	1.71	SEG1
ATCM	562	NZ	LYS	38	4.547	8.742	9.007	1.00	1.03	SEG1
ATCM	563	HZ1	LYS	38	5.129	9.580	9.207	1.00	1.31	SEG1
ATCM	564	HZ2	LYS	38	4.478	8.152	9.862	1.00	1.61	SEG1
ATCM	565	HZ3	LYS	38	3.596	9.044	8.716	1.00	1.33	SEG1
ATCM	566	C	LYS	38	9.619	6.811	5.269	1.00	0.32	SEG1
ATCM	567	O	LYS	38	8.966	6.417	4.321	1.00	0.34	SEG1
ATCM	568	N	VAL	39	10.366	7.883	5.191	1.00	0.33	SEG1
ATCM	569	HN	VAL	39	10.888	8.164	5.969	1.00	0.34	SEG1
ATCM	570	CA	VAL	39	10.439	8.664	3.917	1.00	0.35	SEG1
ATCM	571	HA	VAL	39	9.447	8.907	3.564	1.00	0.36	SEG1
ATCM	572	CB	VAL	39	11.208	9.947	4.274	1.00	0.38	SEG1
ATCM	573	HB	VAL	39	12.149	9.682	4.733	1.00	0.39	SEG1
ATCM	574	CG1	VAL	39	11.479	10.772	3.009	1.00	0.42	SEG1
ATCM	575	HG11	VAL	39	11.881	11.735	3.286	1.00	1.12	SEG1
ATCM	576	HG12	VAL	39	10.557	10.910	2.465	1.00	1.09	SEG1
ATCM	577	HG13	VAL	39	12.190	10.251	2.385	1.00	1.05	SEG1
ATCM	578	CG2	VAL	39	10.381	10.782	5.253	1.00	0.38	SEG1
ATCM	579	HG21	VAL	39	10.683	11.816	5.188	1.00	1.08	SEG1
ATCM	580	HG22	VAL	39	10.545	10.422	6.258	1.00	1.12	SEG1
ATCM	581	HG23	VAL	39	9.334	10.695	5.005	1.00	1.03	SEG1
ATCM	582	C	VAL	39	11.203	7.842	2.870	1.00	0.37	SEG1
ATCM	583	O	VAL	39	10.831	7.796	1.713	1.00	0.40	SEG1
ATCM	584	N	ALA	40	12.260	7.178	3.277	1.00	0.39	SEG1
ATCM	585	HN	ALA	40	12.533	7.222	4.218	1.00	0.38	SEG1
ATCM	586	CA	ALA	40	13.035	6.340	2.310	1.00	0.42	SEG1
ATCM	587	HA	ALA	40	13.366	6.933	1.471	1.00	0.46	SEG1
ATCM	588	CB	ALA	40	14.234	5.810	3.098	1.00	0.45	SEG1
ATCM	589	HB1	ALA	40	14.571	6.566	3.793	1.00	1.04	SEG1
ATCM	590	HB2	ALA	40	15.035	5.566	2.416	1.00	1.12	SEG1
ATCM	591	HB3	ALA	40	13.943	4.924	3.644	1.00	1.11	SEG1
ATCM	592	C	ALA	40	12.142	5.193	1.842	1.00	0.40	SEG1
ATCM	593	O	ALA	40	12.097	4.854	0.675	1.00	0.42	SEG1
ATCM	594	N	VAL	41	11.401	4.625	2.756	1.00	0.38	SEG1
ATCM	595	HN	VAL	41	11.447	4.948	3.681	1.00	0.37	SEG1
ATCM	596	CA	VAL	41	10.457	3.522	2.402	1.00	0.38	SEG1
ATCM	597	HA	VAL	41	10.991	2.697	1.958	1.00	0.41	SEG1
ATCM	598	CB	VAL	41	9.836	3.092	3.741	1.00	0.37	SEG1
ATCM	599	HB	VAL	41	9.493	3.965	4.276	1.00	0.35	SEG1
ATCM	600	CG1	VAL	41	8.663	2.133	3.508	1.00	0.38	SEG1
ATCM	601	HG11	VAL	41	7.767	2.703	3.315	1.00	0.97	SEG1
ATCM	602	HG12	VAL	41	8.518	1.522	4.387	1.00	1.10	SEG1
ATCM	603	HG13	VAL	41	8.878	1.500	2.660	1.00	1.07	SEG1

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ATCM	604	CG2	VAL	41	10.901	2.376	4.573	1.00	0.39	SEG1
ATCM	605	HG21	VAL	41	11.863	2.833	4.395	1.00	1.06	SEG1
ATCM	606	HG22	VAL	41	10.940	1.335	4.290	1.00	1.05	SEG1
ATCM	607	HG23	VAL	41	10.654	2.456	5.622	1.00	1.11	SEG1
ATCM	608	C	VAL	41	9.411	4.070	1.424	1.00	0.37	SEG1
ATCM	609	O	VAL	41	9.028	3.415	0.475	1.00	0.39	SEG1
ATCM	610	N	TYR	42	8.958	5.278	1.655	1.00	0.37	SEG1
ATCM	611	HN	TYR	42	9.295	5.778	2.417	1.00	0.37	SEG1
ATCM	612	CA	TYR	42	7.941	5.909	0.746	1.00	0.40	SEG1
ATCM	613	HA	TYR	42	6.973	5.454	0.886	1.00	0.40	SEG1
ATCM	614	CB	TYR	42	7.896	7.390	1.168	1.00	0.43	SEG1
ATCM	615	HB1	TYR	42	8.104	8.016	0.312	1.00	0.53	SEG1
ATCM	616	HB2	TYR	42	8.641	7.570	1.926	1.00	0.47	SEG1
ATCM	617	CG	TYR	42	6.539	7.749	1.724	1.00	0.41	SEG1
ATCM	618	CD1	TYR	42	5.881	8.890	1.251	1.00	1.26	SEG1
ATCM	619	HD1	TYR	42	6.341	9.499	0.488	1.00	2.09	SEG1
ATCM	620	CD2	TYR	42	5.944	6.960	2.715	1.00	1.17	SEG1
ATCM	621	HD2	TYR	42	6.450	6.081	3.082	1.00	2.02	SEG1
ATCM	622	CE1	TYR	42	4.631	9.242	1.764	1.00	1.31	SEG1
ATCM	623	HE1	TYR	42	4.129	10.125	1.397	1.00	2.16	SEG1
ATCM	624	CE2	TYR	42	4.691	7.311	3.218	1.00	1.19	SEG1
ATCM	625	HE2	TYR	42	4.233	6.703	3.991	1.00	2.02	SEG1
ATCM	626	CZ	TYR	42	4.034	8.453	2.754	1.00	0.60	SEG1
ATCM	627	OH	TYR	42	2.799	8.799	3.261	1.00	0.75	SEG1
ATCM	628	HH	TYR	42	2.917	9.056	4.179	1.00	1.20	SEG1
ATCM	629	C	TYR	42	8.385	5.801	-0.722	1.00	0.43	SEG1
ATCM	630	O	TYR	42	7.621	5.411	-1.584	1.00	0.42	SEG1
ATCM	631	N	ARG	43	9.615	6.155	-1.004	1.00	0.49	SEG1
ATCM	632	HN	ARG	43	10.206	6.471	-0.288	1.00	0.51	SEG1
ATCM	633	CA	ARG	43	10.116	6.086	-2.411	1.00	0.54	SEG1
ATCM	634	HA	ARG	43	9.465	6.640	-3.069	1.00	0.56	SEG1
ATCM	635	CB	ARG	43	11.501	6.736	-2.380	1.00	0.66	SEG1
ATCM	636	HB1	ARG	43	11.994	6.578	-3.327	1.00	1.44	SEG1
ATCM	637	HB2	ARG	43	12.089	6.292	-1.589	1.00	1.20	SEG1
ATCM	638	CG	ARG	43	11.358	8.238	-2.128	1.00	0.88	SEG1
ATCM	639	HG1	ARG	43	10.875	8.398	-1.175	1.00	1.42	SEG1
ATCM	640	HG2	ARG	43	10.762	8.680	-2.914	1.00	1.62	SEG1
ATCM	641	CD	ARG	43	12.746	8.887	-2.107	1.00	0.77	SEG1
ATCM	642	HD1	ARG	43	13.387	8.381	-1.402	1.00	1.04	SEG1
ATCM	643	HD2	ARG	43	12.665	9.937	-1.859	1.00	1.12	SEG1
ATCM	644	NE	ARG	43	13.278	8.717	-3.497	1.00	1.91	SEG1
ATCM	645	HE	ARG	43	12.718	8.291	-4.180	1.00	2.62	SEG1
ATCM	646	CZ	ARG	43	14.490	9.118	-3.823	1.00	2.29	SEG1
ATCM	647	NH1	ARG	43	14.915	8.932	-5.042	1.00	3.44	SEG1
ATCM	648	NH11	ARG	43	14.323	8.491	-5.716	1.00	4.07	SEG1
ATCM	649	NH12	ARG	43	15.832	9.231	-5.303	1.00	3.79	SEG1
ATCM	650	NH2	ARG	43	15.279	9.704	-2.952	1.00	1.85	SEG1
ATCM	651	NH21	ARG	43	14.973	9.860	-2.014	1.00	1.42	SEG1
ATCM	652	NH22	ARG	43	16.195	9.997	-3.230	1.00	2.34	SEG1
ATCM	653	C	ARG	43	10.227	4.630	-2.874	1.00	0.50	SEG1
ATCM	654	O	ARG	43	9.908	4.304	-4.002	1.00	0.50	SEG1
ATCM	655	N	ALA	44	10.680	3.754	-2.011	1.00	0.51	SEG1
ATCM	656	HN	ALA	44	10.931	4.044	-1.108	1.00	0.52	SEG1
ATCM	657	CA	ALA	44	10.816	2.315	-2.400	1.00	0.52	SEG1
ATCM	658	HA	ALA	44	11.458	2.217	-3.261	1.00	0.57	SEG1
ATCM	659	CB	ALA	44	11.454	1.627	-1.191	1.00	0.54	SEG1
ATCM	660	HB1	ALA	44	10.679	1.303	-0.511	1.00	1.08	SEG1
ATCM	661	HB2	ALA	44	12.108	2.321	-0.686	1.00	1.23	SEG1
ATCM	662	HB3	ALA	44	12.023	0.771	-1.522	1.00	1.10	SEG1
ATCM	663	C	ALA	44	9.439	1.714	-2.690	1.00	0.45	SEG1
ATCM	664	O	ALA	44	9.255	0.999	-3.657	1.00	0.47	SEG1
ATCM	665	N	LEU	45	8.474	2.001	-1.855	1.00	0.41	SEG1
ATCM	666	HN	LEU	45	8.655	2.581	-1.084	1.00	0.42	SEG1
ATCM	667	CA	LEU	45	7.101	1.454	-2.069	1.00	0.38	SEG1
ATCM	668	HA	LEU	45	7.139	0.384	-2.197	1.00	0.41	SEG1
ATCM	669	CB	LEU	45	6.323	1.802	-0.801	1.00	0.37	SEG1
ATCM	670	HB1	LEU	45	5.278	1.582	-0.951	1.00	0.37	SEG1
ATCM	671	HB2	LEU	45	6.442	2.854	-0.581	1.00	0.37	SEG1
ATCM	672	CG	LEU	45	6.852	0.971	0.368	1.00	0.40	SEG1
ATCM	673	HG	LEU	45	7.928	1.054	0.410	1.00	0.41	SEG1
ATCM	674	CD1	LEU	45	6.251	1.494	1.673	1.00	0.42	SEG1
ATCM	675	HD11	LEU	45	5.183	1.332	1.658	1.00	0.91	SEG1
ATCM	676	HD12	LEU	45	6.455	2.550	1.764	1.00	1.16	SEG1
ATCM	677	HD13	LEU	45	6.692	0.968	2.507	1.00	1.10	SEG1
ATCM	678	CD2	LEU	45	6.457	-0.495	0.176	1.00	0.45	SEG1
ATCM	679	HD21	LEU	45	7.249	-1.016	-0.341	1.00	1.17	SEG1
ATCM	680	HD22	LEU	45	5.550	-0.551	-0.408	1.00	1.07	SEG1

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ATOM	651	HD23	LEU	45	6.294	-0.954	1.140	1.00	1.06	SEG1
ATOM	652	C	LEU	45	6.457	2.115	-3.290	1.00	0.37	SEG1
ATOM	653	O	LEU	45	5.698	1.500	-4.014	1.00	0.37	SEG1
ATOM	654	N	GLN	46	6.755	3.369	-3.513	1.00	0.37	SEG1
ATOM	655	HN	GLN	46	7.368	3.836	-2.906	1.00	0.39	SEG1
ATOM	656	CA	GLN	46	6.162	4.094	-4.680	1.00	0.38	SEG1
ATOM	657	HA	GLN	46	5.089	4.141	-4.588	1.00	0.39	SEG1
ATOM	658	CB	GLN	46	6.756	5.506	-4.611	1.00	0.42	SEG1
ATOM	659	HB1	GLN	46	7.831	5.447	-4.681	1.00	0.42	SEG1
ATOM	690	HB2	GLN	46	6.482	5.963	-3.671	1.00	0.43	SEG1
ATOM	691	CG	GLN	46	6.220	6.358	-5.767	1.00	0.46	SEG1
ATOM	692	HG1	GLN	46	5.144	6.415	-5.702	1.00	0.54	SEG1
ATOM	693	HG2	GLN	46	6.500	5.905	-6.706	1.00	0.65	SEG1
ATOM	694	CD	GLN	46	6.808	7.772	-5.687	1.00	0.92	SEG1
ATOM	695	OE1	GLN	46	7.395	8.149	-4.690	1.00	1.16	SEG1
ATOM	696	NE2	GLN	46	6.687	8.573	-6.710	1.00	1.46	SEG1
ATOM	697	HE21	GLN	46	6.225	8.269	-7.519	1.00	1.64	SEG1
ATOM	698	HE22	GLN	46	7.057	9.480	-6.669	1.00	1.83	SEG1
ATOM	699	C	GLN	46	6.555	3.402	-5.989	1.00	0.37	SEG1
ATOM	700	O	GLN	46	5.752	3.272	-6.895	1.00	0.37	SEG1
ATOM	701	N	ALA	47	7.781	2.957	-6.092	1.00	0.39	SEG1
ATOM	702	HN	ALA	47	8.405	3.072	-5.345	1.00	0.41	SEG1
ATOM	703	CA	ALA	47	8.226	2.267	-7.341	1.00	0.41	SEG1
ATOM	704	HA	ALA	47	8.001	2.873	-8.204	1.00	0.41	SEG1
ATOM	705	CB	ALA	47	9.740	2.105	-7.195	1.00	0.46	SEG1
ATOM	706	HB1	ALA	47	9.958	1.155	-6.730	1.00	1.14	SEG1
ATOM	707	HB2	ALA	47	10.131	2.904	-6.582	1.00	1.11	SEG1
ATOM	708	HB3	ALA	47	10.202	2.143	-8.171	1.00	1.09	SEG1
ATOM	709	C	ALA	47	7.550	0.898	-7.462	1.00	0.39	SEG1
ATOM	710	O	ALA	47	7.154	0.487	-8.536	1.00	0.39	SEG1
ATOM	711	N	ALA	48	7.422	0.191	-6.368	1.00	0.40	SEG1
ATOM	712	HN	ALA	48	7.755	0.547	-5.516	1.00	0.41	SEG1
ATOM	713	CA	ALA	48	6.777	-1.159	-6.414	1.00	0.42	SEG1
ATOM	714	HA	ALA	48	7.346	-1.823	-7.045	1.00	0.45	SEG1
ATOM	715	CB	ALA	48	6.800	-1.669	-4.972	1.00	0.46	SEG1
ATOM	716	HB1	ALA	48	6.464	-0.887	-4.307	1.00	1.10	SEG1
ATOM	717	HB2	ALA	48	7.807	-1.958	-4.708	1.00	1.01	SEG1
ATOM	718	HB3	ALA	48	6.146	-2.524	-4.882	1.00	0.97	SEG1
ATOM	719	C	ALA	48	5.336	-1.051	-6.921	1.00	0.39	SEG1
ATOM	720	O	ALA	48	4.896	-1.843	-7.734	1.00	0.40	SEG1
ATOM	721	N	LEU	49	4.598	-0.079	-6.446	1.00	0.36	SEG1
ATOM	722	HN	LEU	49	4.977	0.544	-5.790	1.00	0.36	SEG1
ATOM	723	CA	LEU	49	3.181	0.083	-6.896	1.00	0.35	SEG1
ATOM	724	HA	LEU	49	2.603	-0.787	-6.632	1.00	0.38	SEG1
ATOM	725	CB	LEU	49	2.647	1.305	-6.154	1.00	0.33	SEG1
ATOM	726	HB1	LEU	49	1.643	1.509	-6.490	1.00	0.35	SEG1
ATOM	727	HB2	LEU	49	3.275	2.156	-6.362	1.00	0.33	SEG1
ATOM	728	CG	LEU	49	2.634	1.032	-4.650	1.00	0.35	SEG1
ATOM	729	HG	LEU	49	3.604	0.669	-4.341	1.00	0.37	SEG1
ATOM	730	CD1	LEU	49	2.312	2.326	-3.902	1.00	0.37	SEG1
ATOM	731	HD11	LEU	49	2.785	2.309	-2.932	1.00	1.08	SEG1
ATOM	732	HD12	LEU	49	1.242	2.414	-3.781	1.00	1.06	SEG1
ATOM	733	HD13	LEU	49	2.680	3.169	-4.467	1.00	1.10	SEG1
ATOM	734	CD2	LEU	49	1.568	-0.017	-4.329	1.00	0.39	SEG1
ATOM	735	HD21	LEU	49	1.954	-1.002	-4.549	1.00	1.04	SEG1
ATOM	736	HD22	LEU	49	0.689	0.168	-4.928	1.00	1.15	SEG1
ATOM	737	HD23	LEU	49	1.309	0.043	-3.282	1.00	1.02	SEG1
ATOM	738	C	LEU	49	3.124	0.321	-8.406	1.00	0.36	SEG1
ATOM	739	O	LEU	49	2.256	-0.188	-9.089	1.00	0.38	SEG1
ATOM	740	N	ALA	50	4.041	1.101	-8.932	1.00	0.36	SEG1
ATOM	741	HN	ALA	50	4.726	1.504	-8.355	1.00	0.35	SEG1
ATOM	742	CA	ALA	50	4.040	1.388	-10.404	1.00	0.39	SEG1
ATOM	743	HA	ALA	50	3.192	1.998	-10.667	1.00	0.40	SEG1
ATOM	744	CB	ALA	50	5.338	2.158	-10.664	1.00	0.42	SEG1
ATOM	745	HB1	ALA	50	5.640	2.669	-9.761	1.00	1.08	SEG1
ATOM	746	HB2	ALA	50	5.177	2.880	-11.449	1.00	1.10	SEG1
ATOM	747	HB3	ALA	50	6.112	1.468	-10.963	1.00	1.13	SEG1
ATOM	748	C	ALA	50	4.025	0.082	-11.202	1.00	0.41	SEG1
ATOM	749	O	ALA	50	3.347	-0.037	-12.205	1.00	0.44	SEG1
ATOM	750	N	GLU	51	4.759	-0.897	-10.749	1.00	0.42	SEG1
ATOM	751	HN	GLU	51	5.285	-0.771	-9.932	1.00	0.40	SEG1
ATOM	752	CA	GLU	51	4.788	-2.209	-11.460	1.00	0.46	SEG1
ATOM	753	HA	GLU	51	5.054	-2.070	-12.495	1.00	0.50	SEG1
ATOM	754	CB	GLU	51	5.864	-3.033	-10.750	1.00	0.49	SEG1
ATOM	755	HB1	GLU	51	5.823	-4.055	-11.096	1.00	0.53	SEG1
ATOM	756	HB2	GLU	51	5.693	-3.006	-9.684	1.00	0.47	SEG1
ATOM	757	CG	GLU	51	7.245	-2.446	-11.064	1.00	0.52	SEG1

ATCM	758	HG1	GLU	51	7.284	-1.423	-10.720	1.00	0.83	SEG1
ATCM	759	HG2	GLU	51	7.409	-2.470	-12.131	1.00	0.71	SEG1
ATCM	760	CD	GLU	51	8.343	-3.257	-10.361	1.00	0.92	SEG1
ATCM	761	DE1	GLU	51	8.013	-4.095	-9.533	1.00	1.57	SEG1
ATCM	762	DE2	GLU	51	9.502	-3.025	-10.664	1.00	1.58	SEG1
ATCM	763	C	GLU	51	3.422	-2.895	-11.347	1.00	0.46	SEG1
ATCM	764	O	GLU	51	2.956	-3.522	-12.279	1.00	0.50	SEG1
ATCM	765	N	SER	52	2.784	-2.789	-10.205	1.00	0.42	SEG1
ATCM	766	HN	SER	52	3.185	-2.283	-9.466	1.00	0.39	SEG1
ATCM	767	CA	SER	52	1.451	-3.445	-10.022	1.00	0.43	SEG1
ATCM	768	HA	SER	52	1.453	-4.420	-10.480	1.00	0.47	SEG1
ATCM	769	CB	SER	52	1.287	-3.588	-8.510	1.00	0.42	SEG1
ATCM	770	HB1	SER	52	0.312	-4.003	-8.293	1.00	1.06	SEG1
ATCM	771	HB2	SER	52	1.375	-2.622	-8.043	1.00	0.95	SEG1
ATCM	772	OG	SER	52	2.304	-4.445	-8.007	1.00	1.34	SEG1
ATCM	773	HG	SER	52	2.885	-3.922	-7.451	1.00	1.64	SEG1
ATCM	774	C	SER	52	0.316	-2.587	-10.604	1.00	0.42	SEG1
ATCM	775	O	SER	52	-0.808	-3.043	-10.713	1.00	0.46	SEG1
ATCM	776	N	GLY	53	0.591	-1.351	-10.984	1.00	0.41	SEG1
ATCM	777	HN	GLY	53	1.500	-1.009	-10.891	1.00	0.40	SEG1
ATCM	778	CA	GLY	53	-0.483	-0.497	-11.563	1.00	0.42	SEG1
ATCM	779	HA1	GLY	53	-1.350	-1.102	-11.778	1.00	0.45	SEG1
ATCM	780	HA2	GLY	53	-0.125	-0.048	-12.477	1.00	0.45	SEG1
ATCM	781	C	GLY	53	-0.876	0.610	-10.576	1.00	0.39	SEG1
ATCM	782	O	GLY	53	-1.904	1.244	-10.731	1.00	0.41	SEG1
ATCM	783	N	GLY	54	-0.068	0.858	-9.574	1.00	0.37	SEG1
ATCM	784	HN	GLY	54	0.758	0.348	-9.472	1.00	0.38	SEG1
ATCM	785	CA	GLY	54	-0.395	1.933	-8.594	1.00	0.35	SEG1
ATCM	786	HA1	GLY	54	-0.294	1.551	-7.589	1.00	0.35	SEG1
ATCM	787	HA2	GLY	54	-1.408	2.274	-8.754	1.00	0.35	SEG1
ATCM	788	C	GLY	54	0.578	3.091	-8.797	1.00	0.38	SEG1
ATCM	789	O	GLY	54	1.469	3.314	-7.999	1.00	0.40	SEG1
ATCM	790	N	SER	55	0.417	3.823	-9.869	1.00	0.41	SEG1
ATCM	791	HN	SER	55	-0.305	3.610	-10.497	1.00	0.41	SEG1
ATCM	792	CA	SER	55	1.338	4.969	-10.146	1.00	0.46	SEG1
ATCM	793	HA	SER	55	2.352	4.618	-10.258	1.00	0.50	SEG1
ATCM	794	CB	SER	55	0.846	5.576	-11.463	1.00	0.50	SEG1
ATCM	795	HB1	SER	55	0.676	4.784	-12.181	1.00	0.50	SEG1
ATCM	796	HB2	SER	55	1.587	6.254	-11.850	1.00	0.59	SEG1
ATCM	797	OG	SER	55	-0.363	6.286	-11.233	1.00	0.49	SEG1
ATCM	798	HG	SER	55	-0.161	7.225	-11.238	1.00	0.74	SEG1
ATCM	799	C	SER	55	1.248	6.001	-9.012	1.00	0.46	SEG1
ATCM	800	O	SER	55	0.277	6.023	-8.283	1.00	0.42	SEG1
ATCM	801	N	PRO	56	2.265	6.826	-8.889	1.00	0.54	SEG1
ATCM	802	CA	PRO	56	2.267	7.855	-7.818	1.00	0.56	SEG1
ATCM	803	HA	PRO	56	2.081	7.403	-6.858	1.00	0.55	SEG1
ATCM	804	CB	PRO	56	3.685	8.417	-7.861	1.00	0.66	SEG1
ATCM	805	HB1	PRO	56	4.314	7.904	-7.151	1.00	0.69	SEG1
ATCM	806	HB2	PRO	56	3.673	9.480	-7.659	1.00	0.68	SEG1
ATCM	807	CG	PRO	56	4.163	8.154	-9.252	1.00	0.70	SEG1
ATCM	808	HG1	PRO	56	5.233	8.017	-9.258	1.00	0.77	SEG1
ATCM	809	HG2	PRO	56	3.890	8.978	-9.898	1.00	0.72	SEG1
ATCM	810	CD	PRO	56	3.487	6.889	-9.711	1.00	0.64	SEG1
ATCM	811	HD2	PRO	56	3.240	6.952	-10.763	1.00	0.65	SEG1
ATCM	812	HD1	PRO	56	4.111	6.032	-9.515	1.00	0.67	SEG1
ATCM	813	C	PRO	56	1.241	8.958	-8.105	1.00	0.55	SEG1
ATCM	814	O	PRO	56	0.942	9.765	-7.244	1.00	0.57	SEG1
ATCM	815	N	ASP	57	0.695	9.001	-9.298	1.00	0.56	SEG1
ATCM	816	HN	ASP	57	0.940	8.342	-9.978	1.00	0.56	SEG1
ATCM	817	CA	ASP	57	-0.310	10.053	-9.621	1.00	0.60	SEG1
ATCM	818	HA	ASP	57	-0.082	10.966	-9.095	1.00	0.65	SEG1
ATCM	819	CB	ASP	57	-0.185	10.273	-11.129	1.00	0.68	SEG1
ATCM	820	HB1	ASP	57	-0.999	10.894	-11.470	1.00	1.04	SEG1
ATCM	821	HB2	ASP	57	-0.223	9.320	-11.636	1.00	1.22	SEG1
ATCM	822	CG	ASP	57	1.146	10.965	-11.444	1.00	1.17	SEG1
ATCM	823	OD1	ASP	57	1.650	10.762	-12.536	1.00	1.87	SEG1
ATCM	824	OD2	ASP	57	1.636	11.689	-10.591	1.00	1.84	SEG1
ATCM	825	C	ASP	57	-1.713	9.565	-9.264	1.00	0.57	SEG1
ATCM	826	O	ASP	57	-2.603	10.355	-9.006	1.00	0.66	SEG1
ATCM	827	N	VAL	58	-1.921	8.271	-9.259	1.00	0.49	SEG1
ATCM	828	HN	VAL	58	-1.192	7.654	-9.479	1.00	0.47	SEG1
ATCM	829	CA	VAL	58	-3.271	7.737	-8.930	1.00	0.50	SEG1
ATCM	830	HA	VAL	58	-3.985	8.545	-8.933	1.00	0.54	SEG1
ATCM	831	CB	VAL	58	-3.603	6.764	-10.083	1.00	0.51	SEG1
ATCM	832	HB	VAL	58	-3.274	7.198	-11.715	1.00	0.55	SEG1
ATCM	833	CG1	VAL	58	-2.905	5.406	-9.905	1.00	0.46	SEG1
ATCM	834	HG11	VAL	58	-3.317	4.899	-9.045	1.00	1.04	SEG1

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ATOM	835	HG12	VAL	58	-1.849	5.558	-9.760	1.00	1.10	SEG1
ATOM	836	HG13	VAL	58	-3.063	4.803	-10.786	1.00	1.17	SEG1
ATOM	837	CG2	VAL	58	-5.116	6.560	-10.141	1.00	0.64	SEG1
ATOM	838	HG21	VAL	58	-5.442	6.592	-11.169	1.00	1.17	SEG1
ATOM	839	HG22	VAL	58	-5.610	7.344	-9.585	1.00	1.24	SEG1
ATOM	840	HG23	VAL	58	-5.369	5.602	-9.712	1.00	1.13	SEG1
ATOM	841	C	VAL	58	-3.277	7.052	-7.549	1.00	0.45	SEG1
ATOM	842	O	VAL	58	-4.179	7.252	-6.757	1.00	0.49	SEG1
ATOM	843	N	LEU	59	-2.280	6.252	-7.261	1.00	0.40	SEG1
ATOM	844	HN	LEU	59	-1.564	6.110	-7.912	1.00	0.40	SEG1
ATOM	845	CA	LEU	59	-2.224	5.559	-5.937	1.00	0.36	SEG1
ATOM	846	HA	LEU	59	-3.151	5.698	-5.405	1.00	0.38	SEG1
ATOM	847	CB	LEU	59	-2.030	4.076	-6.269	1.00	0.34	SEG1
ATOM	848	HB1	LEU	59	-1.319	3.643	-5.581	1.00	0.32	SEG1
ATOM	849	HB2	LEU	59	-1.659	3.980	-7.279	1.00	0.35	SEG1
ATOM	850	CG	LEU	59	-3.366	3.339	-6.147	1.00	0.35	SEG1
ATOM	851	HG	LEU	59	-4.177	4.025	-6.348	1.00	0.37	SEG1
ATOM	852	CD1	LEU	59	-3.408	2.186	-7.150	1.00	0.36	SEG1
ATOM	853	HD11	LEU	59	-2.581	1.517	-6.964	1.00	1.01	SEG1
ATOM	854	HD12	LEU	59	-3.337	2.579	-8.153	1.00	1.15	SEG1
ATOM	855	HD13	LEU	59	-4.338	1.648	-7.040	1.00	1.03	SEG1
ATOM	856	CD2	LEU	59	-3.506	2.778	-4.730	1.00	0.34	SEG1
ATOM	857	HD21	LEU	59	-3.064	1.793	-4.688	1.00	0.99	SEG1
ATOM	858	HD22	LEU	59	-4.552	2.714	-4.469	1.00	0.92	SEG1
ATOM	859	HD23	LEU	59	-3.000	3.430	-4.033	1.00	1.10	SEG1
ATOM	860	C	LEU	59	-1.053	6.088	-5.106	1.00	0.34	SEG1
ATOM	861	O	LEU	59	0.087	6.060	-5.532	1.00	0.36	SEG1
ATOM	862	N	GLN	60	-1.331	6.565	-3.921	1.00	0.34	SEG1
ATOM	863	HN	GLN	60	-2.259	6.572	-3.605	1.00	0.36	SEG1
ATOM	864	CA	GLN	60	-0.248	7.099	-3.044	1.00	0.34	SEG1
ATOM	865	HA	GLN	60	0.721	6.849	-3.443	1.00	0.35	SEG1
ATOM	866	CB	GLN	60	-0.447	8.615	-3.053	1.00	0.39	SEG1
ATOM	867	HB1	GLN	60	0.187	9.067	-2.304	1.00	0.61	SEG1
ATOM	868	HB2	GLN	60	-1.480	8.843	-2.835	1.00	0.60	SEG1
ATOM	869	CG	GLN	60	-0.081	9.170	-4.430	1.00	0.80	SEG1
ATOM	870	HG1	GLN	60	-0.599	8.610	-5.194	1.00	1.24	SEG1
ATOM	871	HG2	GLN	60	0.986	9.084	-4.581	1.00	1.15	SEG1
ATOM	872	CD	GLN	60	-0.492	10.641	-4.513	1.00	1.13	SEG1
ATOM	873	OE1	GLN	60	-1.398	11.070	-3.827	1.00	1.43	SEG1
ATOM	874	NE2	GLN	60	0.141	11.438	-5.330	1.00	1.63	SEG1
ATOM	875	HE21	GLN	60	0.873	11.093	-5.882	1.00	1.87	SEG1
ATOM	876	HE22	GLN	60	-0.115	12.382	-5.390	1.00	1.98	SEG1
ATOM	877	C	GLN	60	-0.406	6.545	-1.626	1.00	0.30	SEG1
ATOM	878	O	GLN	60	-1.482	6.135	-1.231	1.00	0.30	SEG1
ATOM	879	N	MET	61	0.654	6.535	-0.857	1.00	0.30	SEG1
ATOM	880	HN	MET	61	1.508	6.874	-1.196	1.00	0.33	SEG1
ATOM	881	CA	MET	61	0.560	6.012	0.539	1.00	0.28	SEG1
ATOM	882	HA	MET	61	-0.211	5.264	0.609	1.00	0.27	SEG1
ATOM	883	CB	MET	61	1.926	5.387	0.820	1.00	0.31	SEG1
ATOM	884	HB1	MET	61	1.999	5.139	1.868	1.00	0.35	SEG1
ATOM	885	HB2	MET	61	2.704	6.090	0.560	1.00	0.35	SEG1
ATOM	886	CG	MET	61	2.084	4.116	-0.014	1.00	0.32	SEG1
ATOM	887	HG1	MET	61	1.880	4.339	-1.051	1.00	0.64	SEG1
ATOM	888	HG2	MET	61	1.389	3.368	0.337	1.00	0.57	SEG1
ATOM	889	SD	MET	61	3.773	3.494	0.150	1.00	0.71	SEG1
ATOM	890	CE	MET	61	4.507	4.495	-1.167	1.00	0.46	SEG1
ATOM	891	HE1	MET	61	5.558	4.643	-0.963	1.00	1.11	SEG1
ATOM	892	HE2	MET	61	4.013	5.453	-1.210	1.00	1.09	SEG1
ATOM	893	HE3	MET	61	4.385	3.987	-2.114	1.00	1.09	SEG1
ATOM	894	C	MET	61	0.284	7.164	1.505	1.00	0.30	SEG1
ATOM	895	O	MET	61	1.066	8.090	1.617	1.00	0.40	SEG1
ATOM	896	N	LEU	62	-0.829	7.119	2.191	1.00	0.29	SEG1
ATOM	897	HN	LEU	62	-1.445	6.365	2.073	1.00	0.35	SEG1
ATOM	898	CA	LEU	62	-1.168	8.220	3.144	1.00	0.32	SEG1
ATOM	899	HA	LEU	62	-1.099	9.175	2.645	1.00	0.36	SEG1
ATOM	900	CB	LEU	62	-2.618	7.961	3.556	1.00	0.33	SEG1
ATOM	901	HB1	LEU	62	-2.884	8.613	4.373	1.00	0.84	SEG1
ATOM	902	HB2	LEU	62	-2.727	6.930	3.865	1.00	0.78	SEG1
ATOM	903	CG	LEU	62	-3.536	8.239	2.364	1.00	1.02	SEG1
ATOM	904	HG	LEU	62	-3.166	7.713	1.496	1.00	1.76	SEG1
ATOM	905	CD1	LEU	62	-4.952	7.760	2.683	1.00	1.40	SEG1
ATOM	906	HD11	LEU	62	-4.962	6.682	2.744	1.00	1.72	SEG1
ATOM	907	HD12	LEU	62	-5.625	8.082	1.902	1.00	2.05	SEG1
ATOM	908	HD13	LEU	62	-5.270	3.178	3.626	1.00	1.83	SEG1
ATOM	909	CD2	LEU	62	-3.557	9.742	2.080	1.00	1.50	SEG1
ATOM	910	HD21	LEU	62	-4.406	9.980	1.456	1.00	2.18	SEG1
ATOM	911	HD22	LEU	62	-2.647	10.025	1.572	1.00	1.77	SEG1

ATOM	912	HD23	LEU	62	-3.632	10.284	3.011	1.00	1.93	SEG1
ATOM	913	C	LEU	62	-0.242	8.195	4.363	1.00	0.31	SEG1
ATOM	914	O	LEU	62	0.188	9.230	4.837	1.00	0.34	SEG1
ATOM	915	N	LYS	63	0.076	7.026	4.974	1.00	0.31	SEG1
ATOM	916	HN	LYS	63	-0.278	6.202	4.477	1.00	0.34	SEG1
ATOM	917	CA	LYS	63	0.984	6.956	6.062	1.00	0.33	SEG1
ATOM	918	HA	LYS	63	1.847	7.582	5.928	1.00	0.37	SEG1
ATOM	919	CB	LYS	63	0.154	7.496	7.235	1.00	0.37	SEG1
ATOM	920	HB1	LYS	63	-0.723	6.880	7.368	1.00	0.67	SEG1
ATOM	921	HB2	LYS	63	-0.150	8.511	7.023	1.00	0.76	SEG1
ATOM	922	CG	LYS	63	0.990	7.471	8.522	1.00	0.78	SEG1
ATOM	923	HG1	LYS	63	1.867	8.087	8.393	1.00	1.25	SEG1
ATOM	924	HG2	LYS	63	1.291	6.456	8.735	1.00	1.17	SEG1
ATOM	925	CD	LYS	63	0.157	8.011	9.666	1.00	0.83	SEG1
ATOM	926	HD1	LYS	63	-0.722	7.399	9.815	1.00	1.08	SEG1
ATOM	927	HD2	LYS	63	-0.139	9.026	9.471	1.00	1.00	SEG1
ATOM	928	CE	LYS	63	0.990	7.984	10.970	1.00	1.32	SEG1
ATOM	929	HE1	LYS	63	1.895	8.558	10.846	1.00	1.73	SEG1
ATOM	930	HE2	LYS	63	1.223	6.965	11.244	1.00	1.74	SEG1
ATOM	931	NZ	LYS	63	0.123	8.617	12.004	1.00	1.53	SEG1
ATOM	932	H21	LYS	63	-0.025	9.618	11.768	1.00	1.85	SEG1
ATOM	933	H22	LYS	63	0.583	8.544	12.934	1.00	1.93	SEG1
ATOM	934	H23	LYS	63	-0.796	8.130	12.032	1.00	1.91	SEG1
ATOM	935	C	LYS	63	1.419	5.514	6.336	1.00	0.29	SEG1
ATOM	936	O	LYS	63	0.778	4.566	5.924	1.00	0.28	SEG1
ATOM	937	N	ILE	64	2.499	5.357	7.056	1.00	0.28	SEG1
ATOM	938	HN	ILE	64	2.978	6.141	7.385	1.00	0.31	SEG1
ATOM	939	CA	ILE	64	2.990	3.993	7.402	1.00	0.27	SEG1
ATOM	940	HA	ILE	64	2.589	3.257	6.724	1.00	0.26	SEG1
ATOM	941	CB	ILE	64	4.521	4.064	7.279	1.00	0.29	SEG1
ATOM	942	HB	ILE	64	4.905	4.737	8.033	1.00	0.30	SEG1
ATOM	943	CG1	ILE	64	4.931	4.574	5.883	1.00	0.31	SEG1
ATOM	944	HG11	ILE	64	6.011	4.594	5.813	1.00	0.33	SEG1
ATOM	945	HG12	ILE	64	4.547	5.574	5.742	1.00	0.32	SEG1
ATOM	946	CG2	ILE	64	5.114	2.671	7.507	1.00	0.30	SEG1
ATOM	947	HG21	ILE	64	6.108	2.629	7.085	1.00	1.02	SEG1
ATOM	948	HG22	ILE	64	4.489	1.931	7.030	1.00	1.02	SEG1
ATOM	949	HG23	ILE	64	5.165	2.471	8.567	1.00	1.04	SEG1
ATOM	950	CD1	ILE	64	4.368	3.654	4.791	1.00	0.30	SEG1
ATOM	951	HD11	ILE	64	4.830	3.894	3.844	1.00	1.02	SEG1
ATOM	952	HD12	ILE	64	3.300	3.796	4.714	1.00	1.05	SEG1
ATOM	953	HD13	ILE	64	4.577	2.625	5.042	1.00	1.09	SEG1
ATOM	954	C	ILE	64	2.583	3.686	8.847	1.00	0.26	SEG1
ATOM	955	O	ILE	64	3.032	4.337	9.773	1.00	0.29	SEG1
ATOM	956	N	HIS	65	1.728	2.716	9.047	1.00	0.27	SEG1
ATOM	957	HN	HIS	65	1.371	2.214	8.284	1.00	0.28	SEG1
ATOM	958	CA	HIS	65	1.280	2.384	10.435	1.00	0.27	SEG1
ATOM	959	HA	HIS	65	1.387	3.243	11.077	1.00	0.30	SEG1
ATOM	960	CB	HIS	65	-0.200	2.025	10.296	1.00	0.28	SEG1
ATOM	961	HB1	HIS	65	-0.543	1.551	11.203	1.00	0.31	SEG1
ATOM	962	HB2	HIS	65	-0.329	1.350	9.464	1.00	0.29	SEG1
ATOM	963	CG	HIS	65	-0.999	3.275	10.051	1.00	0.30	SEG1
ATOM	964	ND1	HIS	65	-1.426	4.092	11.086	1.00	1.11	SEG1
ATOM	965	HD1	HIS	65	-1.262	3.949	12.041	1.00	1.90	SEG1
ATOM	966	CD2	HIS	65	-1.456	3.862	8.897	1.00	0.97	SEG1
ATOM	967	HD2	HIS	65	-1.297	3.479	7.900	1.00	1.89	SEG1
ATOM	968	CE1	HIS	65	-2.107	5.115	10.540	1.00	0.85	SEG1
ATOM	969	HE1	HIS	65	-2.560	5.914	11.109	1.00	1.46	SEG1
ATOM	970	NE2	HIS	65	-2.156	5.024	9.208	1.00	0.58	SEG1
ATOM	971	C	HIS	65	2.072	1.201	10.995	1.00	0.28	SEG1
ATOM	972	O	HIS	65	2.183	0.163	10.372	1.00	0.27	SEG1
ATOM	973	N	ARG	66	2.628	1.359	12.170	1.00	0.35	SEG1
ATOM	974	HN	ARG	66	2.527	2.208	12.647	1.00	0.41	SEG1
ATOM	975	CA	ARG	66	3.420	0.258	12.786	1.00	0.39	SEG1
ATOM	976	HA	ARG	66	3.636	-0.510	12.058	1.00	0.36	SEG1
ATOM	977	CB	ARG	66	4.712	0.930	13.235	1.00	0.49	SEG1
ATOM	978	HB1	ARG	66	5.293	0.239	13.821	1.00	0.54	SEG1
ATOM	979	HB2	ARG	66	4.477	1.800	13.831	1.00	0.51	SEG1
ATOM	980	CG	ARG	66	5.516	1.354	12.005	1.00	0.54	SEG1
ATOM	981	HG1	ARG	66	4.935	2.048	11.417	1.00	0.61	SEG1
ATOM	982	HG2	ARG	66	5.748	0.483	11.410	1.00	0.71	SEG1
ATOM	983	CD	ARG	66	6.815	2.029	12.448	1.00	0.64	SEG1
ATOM	984	HD1	ARG	66	7.385	2.345	11.592	1.00	1.19	SEG1
ATOM	985	HD2	ARG	66	7.396	1.353	13.060	1.00	0.78	SEG1
ATOM	986	NE	ARG	66	6.385	3.219	13.248	1.00	1.37	SEG1
ATOM	987	HE	ARG	66	6.366	3.156	14.225	1.00	1.94	SEG1
ATOM	988	CZ	ARG	66	6.041	4.354	12.669	1.00	2.05	SEG1

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ATOM	989	NH1	ARG	66	5.696	5.367	13.415	1.00	2.83	SEG1
ATOM	990	HH11	ARG	66	5.693	5.277	14.411	1.00	3.08	SEG1
ATOM	991	HH12	ARG	66	5.435	6.234	12.991	1.00	3.45	SEG1
ATOM	992	NH2	ARG	66	6.027	4.487	11.362	1.00	2.58	SEG1
ATOM	993	HH21	ARG	66	6.277	3.723	10.771	1.00	2.48	SEG1
ATOM	994	HH22	ARG	66	5.764	5.362	10.956	1.00	3.40	SEG1
ATOM	995	C	ARG	66	2.674	-0.330	13.990	1.00	0.43	SEG1
ATOM	996	O	ARG	66	2.379	0.364	14.945	1.00	0.49	SEG1
ATOM	997	N	SER	67	2.379	-1.604	13.954	1.00	0.42	SEG1
ATOM	998	HN	SER	67	2.635	-2.141	13.175	1.00	0.39	SEG1
ATOM	999	CA	SER	67	1.662	-2.250	15.097	1.00	0.49	SEG1
ATOM	1000	HA	SER	67	1.880	-1.735	16.019	1.00	0.54	SEG1
ATOM	1001	CB	SER	67	0.177	-2.121	14.760	1.00	0.52	SEG1
ATOM	1002	HB1	SER	67	-0.042	-2.702	13.874	1.00	0.53	SEG1
ATOM	1003	HB2	SER	67	-0.064	-1.087	14.576	1.00	0.53	SEG1
ATOM	1004	OG	SER	67	-0.596	-2.593	15.856	1.00	0.57	SEG1
ATOM	1005	HG	SER	67	-0.742	-3.534	15.732	1.00	1.01	SEG1
ATOM	1006	C	SER	67	2.070	-3.721	15.193	1.00	0.49	SEG1
ATOM	1007	O	SER	67	2.287	-4.364	14.187	1.00	0.47	SEG1
ATOM	1008	N	ASP	68	2.182	-4.252	16.398	1.00	0.54	SEG1
ATOM	1009	HN	ASP	68	2.003	-3.695	17.185	1.00	0.56	SEG1
ATOM	1010	CA	ASP	68	2.584	-5.694	16.591	1.00	0.57	SEG1
ATOM	1011	HA	ASP	68	2.852	-5.851	17.625	1.00	0.61	SEG1
ATOM	1012	CB	ASP	68	1.298	-6.497	16.270	1.00	0.60	SEG1
ATOM	1013	HB1	ASP	68	0.449	-5.962	16.671	1.00	0.62	SEG1
ATOM	1014	HB2	ASP	68	1.349	-7.461	16.746	1.00	0.64	SEG1
ATOM	1015	CG	ASP	68	1.092	-6.685	14.758	1.00	0.56	SEG1
ATOM	1016	OD1	ASP	68	1.946	-7.284	14.125	1.00	1.22	SEG1
ATOM	1017	OD2	ASP	68	0.080	-6.221	14.260	1.00	1.18	SEG1
ATOM	1018	C	ASP	68	3.799	-6.044	15.691	1.00	0.53	SEG1
ATOM	1019	O	ASP	68	4.412	-5.146	15.145	1.00	0.49	SEG1
ATOM	1020	N	PRO	69	4.131	-7.314	15.552	1.00	0.55	SEG1
ATOM	1021	CA	PRO	69	5.288	-7.686	14.697	1.00	0.54	SEG1
ATOM	1022	HA	PRO	69	6.116	-7.018	14.871	1.00	0.53	SEG1
ATOM	1023	CB	PRO	69	5.649	-9.089	15.170	1.00	0.60	SEG1
ATOM	1024	HB1	PRO	69	6.420	-9.043	15.922	1.00	0.63	SEG1
ATOM	1025	HB2	PRO	69	5.974	-9.691	14.333	1.00	0.60	SEG1
ATOM	1026	CG	PRO	69	4.393	-9.652	15.760	1.00	0.64	SEG1
ATOM	1027	HG1	PRO	69	4.631	-10.242	16.631	1.00	0.69	SEG1
ATOM	1028	HG2	PRO	69	3.889	-10.267	15.027	1.00	0.64	SEG1
ATOM	1029	CD	PRO	69	3.503	-8.498	16.160	1.00	0.62	SEG1
ATOM	1030	HD2	PRO	69	2.509	-8.648	15.769	1.00	0.62	SEG1
ATOM	1031	HD1	PRO	69	3.477	-8.395	17.234	1.00	0.67	SEG1
ATOM	1032	C	PRO	69	4.905	-7.692	13.206	1.00	0.51	SEG1
ATOM	1033	O	PRO	69	5.179	-8.639	12.494	1.00	0.71	SEG1
ATOM	1034	N	GLN	70	4.290	-6.636	12.729	1.00	0.39	SEG1
ATOM	1035	HN	GLN	70	4.095	-5.881	13.315	1.00	0.48	SEG1
ATOM	1036	CA	GLN	70	3.901	-6.556	11.289	1.00	0.34	SEG1
ATOM	1037	HA	GLN	70	4.557	-7.162	10.678	1.00	0.34	SEG1
ATOM	1038	CB	GLN	70	2.465	-7.086	11.214	1.00	0.37	SEG1
ATOM	1039	HB1	GLN	70	2.064	-6.894	10.230	1.00	0.35	SEG1
ATOM	1040	HB2	GLN	70	1.859	-6.583	11.953	1.00	0.40	SEG1
ATOM	1041	CG	GLN	70	2.446	-8.593	11.481	1.00	0.42	SEG1
ATOM	1042	HG1	GLN	70	2.834	-8.790	12.467	1.00	0.45	SEG1
ATOM	1043	HG2	GLN	70	3.054	-9.098	10.747	1.00	0.41	SEG1
ATOM	1044	CD	GLN	70	1.008	-9.109	11.399	1.00	0.47	SEG1
ATOM	1045	OE1	GLN	70	0.070	-8.372	11.633	1.00	0.63	SEG1
ATOM	1046	NE2	GLN	70	0.794	-10.353	11.070	1.00	0.47	SEG1
ATOM	1047	HE21	GLN	70	1.552	-10.946	10.880	1.00	0.53	SEG1
ATOM	1048	HE22	GLN	70	-0.122	-10.695	11.011	1.00	0.51	SEG1
ATOM	1049	C	GLN	70	3.944	-5.090	10.849	1.00	0.31	SEG1
ATOM	1050	O	GLN	70	3.883	-4.195	11.671	1.00	0.36	SEG1
ATOM	1051	N	LEU	71	4.042	-4.831	9.570	1.00	0.28	SEG1
ATOM	1052	HN	LEU	71	4.087	-5.567	8.921	1.00	0.31	SEG1
ATOM	1053	CA	LEU	71	4.081	-3.413	9.097	1.00	0.26	SEG1
ATOM	1054	HA	LEU	71	4.186	-2.741	9.933	1.00	0.25	SEG1
ATOM	1055	CB	LEU	71	5.316	-3.329	8.200	1.00	0.27	SEG1
ATOM	1056	HB1	LEU	71	5.213	-4.021	7.378	1.00	0.29	SEG1
ATOM	1057	HB2	LEU	71	6.196	-3.582	8.776	1.00	0.28	SEG1
ATOM	1058	CG	LEU	71	5.462	-1.908	7.650	1.00	0.27	SEG1
ATOM	1059	HG	LEU	71	4.550	-1.621	7.146	1.00	0.28	SEG1
ATOM	1060	CD1	LEU	71	5.737	-0.936	8.801	1.00	0.28	SEG1
ATOM	1061	HD11	LEU	71	4.801	-0.569	9.194	1.00	1.03	SEG1
ATOM	1062	HD12	LEU	71	6.325	-0.106	8.437	1.00	1.04	SEG1
ATOM	1063	HD13	LEU	71	6.280	-1.447	9.581	1.00	1.09	SEG1
ATOM	1064	CD2	LEU	71	6.628	-1.868	6.663	1.00	0.31	SEG1
ATOM	1065	HD21	LEU	71	6.253	-1.997	5.659	1.00	1.05	SEG1

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ATOM	1066	HD22	LEU	71	7.320	-2.652	6.894	1.00	1.05	SEG1
ATOM	1067	HD23	LEU	71	7.132	-0.917	6.740	1.00	1.09	SEG1
ATOM	1068	C	LEU	71	2.806	-3.091	8.310	1.00	0.25	SEG1
ATOM	1069	N	LEU	71	2.418	-3.825	7.421	1.00	0.27	SEG1
ATOM	1070	N	ILE	72	2.149	-2.005	8.639	1.00	0.23	SEG1
ATOM	1071	HN	ILE	72	2.478	-1.433	9.362	1.00	0.23	SEG1
ATOM	1072	CA	ILE	72	0.892	-1.642	9.917	1.00	0.23	SEG1
ATOM	1073	HA	ILE	72	0.559	-2.464	7.305	1.00	0.25	SEG1
ATOM	1074	CB	ILE	72	-0.139	-1.356	9.017	1.00	0.24	SEG1
ATOM	1075	HB	ILE	72	0.210	-0.540	9.631	1.00	0.24	SEG1
ATOM	1076	CG1	ILE	72	-0.335	-2.600	9.886	1.00	0.27	SEG1
ATOM	1077	HG11	ILE	72	0.624	-2.961	10.222	1.00	0.27	SEG1
ATOM	1078	HG12	ILE	72	-0.829	-3.364	9.312	1.00	0.30	SEG1
ATOM	1079	CG2	ILE	72	-1.481	-0.977	8.384	1.00	0.26	SEG1
ATOM	1080	HG21	ILE	72	-1.631	-1.553	7.483	1.00	1.07	SEG1
ATOM	1081	HG22	ILE	72	-1.481	0.076	8.143	1.00	1.08	SEG1
ATOM	1082	HG23	ILE	72	-2.279	-1.186	9.081	1.00	0.93	SEG1
ATOM	1083	CD1	ILE	72	-1.194	-2.239	11.097	1.00	0.30	SEG1
ATOM	1084	HD11	ILE	72	-0.584	-1.745	11.838	1.00	1.01	SEG1
ATOM	1085	HD12	ILE	72	-1.617	-3.139	11.519	1.00	1.05	SEG1
ATOM	1086	HD13	ILE	72	-1.991	-1.578	10.786	1.00	1.01	SEG1
ATOM	1087	C	ILE	72	1.114	-0.390	7.061	1.00	0.22	SEG1
ATOM	1088	O	ILE	72	1.602	0.618	7.535	1.00	0.22	SEG1
ATOM	1089	N	VAL	73	0.737	-0.446	5.811	1.00	0.23	SEG1
ATOM	1090	HN	VAL	73	0.333	-1.269	5.462	1.00	0.24	SEG1
ATOM	1091	CA	VAL	73	0.895	0.741	4.916	1.00	0.23	SEG1
ATOM	1092	HA	VAL	73	1.357	1.557	5.450	1.00	0.24	SEG1
ATOM	1093	CB	VAL	73	1.799	0.273	3.771	1.00	0.26	SEG1
ATOM	1094	HB	VAL	73	1.328	-0.551	3.255	1.00	1.09	SEG1
ATOM	1095	CG1	VAL	73	2.023	1.427	2.790	1.00	1.38	SEG1
ATOM	1096	HG11	VAL	73	2.971	1.295	2.290	1.00	1.97	SEG1
ATOM	1097	HG12	VAL	73	2.027	2.363	3.329	1.00	2.06	SEG1
ATOM	1098	HG13	VAL	73	1.229	1.436	2.059	1.00	1.87	SEG1
ATOM	1099	CG2	VAL	73	3.150	-0.178	4.334	1.00	1.35	SEG1
ATOM	1100	HG21	VAL	73	3.022	-1.098	4.885	1.00	1.90	SEG1
ATOM	1101	HG22	VAL	73	3.540	0.585	4.991	1.00	1.99	SEG1
ATOM	1102	HG23	VAL	73	3.842	-0.340	3.521	1.00	1.95	SEG1
ATOM	1103	C	VAL	73	-0.483	1.152	4.386	1.00	0.22	SEG1
ATOM	1104	O	VAL	73	-1.230	0.325	3.896	1.00	0.23	SEG1
ATOM	1105	N	GLN	74	-0.833	2.412	4.485	1.00	0.22	SEG1
ATOM	1106	HN	GLN	74	-0.219	3.066	4.891	1.00	0.24	SEG1
ATOM	1107	CA	GLN	74	-2.175	2.849	3.989	1.00	0.22	SEG1
ATOM	1108	HA	GLN	74	-2.902	2.065	4.130	1.00	0.22	SEG1
ATOM	1109	CB	GLN	74	-2.544	4.062	4.843	1.00	0.25	SEG1
ATOM	1110	HB1	GLN	74	-1.839	4.859	4.658	1.00	0.26	SEG1
ATOM	1111	HB2	GLN	74	-2.516	3.793	5.888	1.00	0.26	SEG1
ATOM	1112	CG	GLN	74	-3.952	4.536	4.468	1.00	0.28	SEG1
ATOM	1113	HG1	GLN	74	-4.634	3.699	4.510	1.00	0.28	SEG1
ATOM	1114	HG2	GLN	74	-3.940	4.938	3.466	1.00	0.28	SEG1
ATOM	1115	CD	GLN	74	-4.419	5.618	5.447	1.00	0.35	SEG1
ATOM	1116	OE1	GLN	74	-3.790	5.858	6.459	1.00	0.45	SEG1
ATOM	1117	NE2	GLN	74	-5.513	6.281	5.189	1.00	0.49	SEG1
ATOM	1118	HE21	GLN	74	-6.026	6.083	4.377	1.00	0.62	SEG1
ATOM	1119	HE22	GLN	74	-5.823	6.976	5.806	1.00	0.55	SEG1
ATOM	1120	C	GLN	74	-2.091	3.243	2.513	1.00	0.23	SEG1
ATOM	1121	O	GLN	74	-1.270	4.053	2.130	1.00	0.26	SEG1
ATOM	1122	N	LEU	75	-2.936	2.678	1.683	1.00	0.22	SEG1
ATOM	1123	HN	LEU	75	-3.589	2.029	2.018	1.00	0.22	SEG1
ATOM	1124	CA	LEU	75	-2.905	3.023	0.229	1.00	0.24	SEG1
ATOM	1125	HA	LEU	75	-2.012	3.580	-0.002	1.00	0.27	SEG1
ATOM	1126	CB	LEU	75	-2.884	1.681	-0.516	1.00	0.24	SEG1
ATOM	1127	HB1	LEU	75	-2.825	1.861	-1.580	1.00	0.30	SEG1
ATOM	1128	HB2	LEU	75	-3.792	1.138	-0.296	1.00	0.28	SEG1
ATOM	1129	CG	LEU	75	-1.673	0.841	-0.080	1.00	0.27	SEG1
ATOM	1130	HG	LEU	75	-1.797	0.552	0.954	1.00	0.62	SEG1
ATOM	1131	CD1	LEU	75	-1.597	-0.414	-0.948	1.00	0.68	SEG1
ATOM	1132	HD11	LEU	75	-1.949	-0.184	-1.943	1.00	1.14	SEG1
ATOM	1133	HD12	LEU	75	-2.214	-1.187	-0.517	1.00	1.42	SEG1
ATOM	1134	HD13	LEU	75	-0.574	-0.755	-0.998	1.00	1.28	SEG1
ATOM	1135	CD2	LEU	75	-0.372	1.642	-0.241	1.00	0.52	SFG1
ATOM	1136	HD21	LEU	75	-0.490	2.365	-1.035	1.00	1.19	SEG1
ATOM	1137	HD22	LEU	75	0.437	0.970	-0.484	1.00	1.30	SEG1
ATOM	1138	HD23	LEU	75	-0.147	2.154	0.684	1.00	1.09	SEG1
ATOM	1139	C	LEU	75	-4.152	3.830	-0.150	1.00	0.25	SEG1
ATOM	1140	O	LEU	75	-5.206	3.672	0.436	1.00	0.26	SEG1
ATOM	1141	N	ARG	76	-4.034	4.693	-1.127	1.00	0.29	SEG1
ATOM	1142	HN	ARG	76	-3.171	4.798	-1.580	1.00	0.32	SEG1

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ATOM	1143	CA	AFG	76	-5.203	5.520	-1.557	1.00	0.32	SEG1
ATOM	1144	HA	AFG	76	-6.060	5.318	-0.935	1.00	0.32	SEG1
ATOM	1145	CB	AFG	76	-4.748	6.970	-1.372	1.00	0.37	SEG1
ATOM	1146	HB1	AFG	76	-3.882	7.157	-1.989	1.00	0.64	SEG1
ATOM	1147	HB2	AFG	76	-4.494	7.139	-0.335	1.00	0.71	SEG1
ATOM	1148	CG	AFG	76	-5.876	7.920	-1.791	1.00	0.83	SEG1
ATOM	1149	HG1	AFG	76	-6.743	7.735	-1.164	1.00	1.20	SEG1
ATOM	1150	HG3	AFG	76	-6.130	7.750	-2.817	1.00	1.15	SEG1
ATOM	1151	HD	AFG	76	-5.423	9.370	-1.597	1.00	0.84	SEG1
ATOM	1152	HD1	AFG	76	-4.576	9.583	-2.230	1.00	1.07	SEG1
ATOM	1153	HD2	AFG	76	-5.177	9.558	-0.560	1.00	1.13	SEG1
ATOM	1154	HE	AFG	76	-6.597	10.197	-2.018	1.00	1.21	SEG1
ATOM	1155	HE	AFG	76	-7.484	9.782	-2.069	1.00	1.76	SEG1
ATOM	1156	HZ	AFG	76	-6.467	11.473	-2.321	1.00	1.51	SEG1
ATOM	1157	NH1	AFG	76	-7.523	12.150	-2.683	1.00	2.15	SEG1
ATOM	1158	NH11	AFG	76	-8.415	11.700	-2.726	1.00	2.53	SEG1
ATOM	1159	NH12	AFG	76	-7.440	13.119	-2.917	1.00	2.50	SEG1
ATOM	1160	NH2	AFG	76	-5.302	12.078	-2.268	1.00	1.85	SEG1
ATOM	1161	NH21	AFG	76	-4.483	11.576	-1.994	1.00	1.91	SEG1
ATOM	1162	NH22	AFG	76	-5.235	13.048	-2.505	1.00	2.40	SEG1
ATOM	1163	C	AFG	76	-5.526	5.241	-3.029	1.00	0.32	SEG1
ATOM	1164	C	AFG	76	-4.637	5.099	-3.845	1.00	0.37	SEG1
ATOM	1165	N	PHE	77	-6.790	5.165	-3.376	1.00	0.32	SEG1
ATOM	1166	HN	PHE	77	-7.492	5.289	-2.701	1.00	0.34	SEG1
ATOM	1167	CA	PHE	77	-7.160	4.895	-4.800	1.00	0.34	SEG1
ATOM	1168	HA	PHE	77	-6.278	4.660	-5.377	1.00	0.34	SEG1
ATOM	1169	CB	PHE	77	-8.093	3.681	-4.759	1.00	0.35	SEG1
ATOM	1170	HB1	PHE	77	-8.486	3.502	-5.746	1.00	0.38	SEG1
ATOM	1171	HB2	PHE	77	-8.908	3.872	-4.076	1.00	0.38	SEG1
ATOM	1172	CG	PHE	77	-7.324	2.467	-4.308	1.00	0.34	SEG1
ATOM	1173	CD1	PHE	77	-6.691	1.653	-5.253	1.00	1.33	SEG1
ATOM	1174	HD1	PHE	77	-6.750	1.900	-6.303	1.00	2.22	SEG1
ATOM	1175	HD2	PHE	77	-7.248	2.149	-2.946	1.00	1.14	SEG1
ATOM	1176	HD2	PHE	77	-7.737	2.778	-2.216	1.00	2.04	SEG1
ATOM	1177	CE1	PHE	77	-5.981	0.521	-4.839	1.00	1.37	SEG1
ATOM	1178	HE1	PHE	77	-5.493	-0.107	-5.570	1.00	2.28	SEG1
ATOM	1179	CE2	PHE	77	-6.537	1.017	-2.531	1.00	1.13	SEG1
ATOM	1180	HE2	PHE	77	-6.478	0.772	-1.480	1.00	2.01	SEG1
ATOM	1181	CZ	PHE	77	-5.904	0.203	-3.477	1.00	0.45	SEG1
ATOM	1182	HZ	PHE	77	-5.356	-0.671	-3.157	1.00	0.53	SEG1
ATOM	1183	C	PHE	77	-7.881	6.105	-5.399	1.00	0.41	SEG1
ATOM	1184	O	PHE	77	-8.479	6.894	-4.693	1.00	0.45	SEG1
ATOM	1185	N	CYS	78	-7.815	6.257	-6.696	1.00	0.48	SEG1
ATOM	1186	HN	CYS	78	-7.319	5.608	-7.233	1.00	0.52	SEG1
ATOM	1187	CA	CYS	78	-8.484	7.423	-7.356	1.00	0.55	SEG1
ATOM	1188	HA	CYS	78	-9.355	7.722	-6.795	1.00	0.58	SEG1
ATOM	1189	CB	CYS	78	-7.438	8.538	-7.320	1.00	0.65	SEG1
ATOM	1190	HB1	CYS	78	-6.626	8.291	-7.989	1.00	0.80	SEG1
ATOM	1191	HB2	CYS	78	-7.056	8.641	-6.315	1.00	0.92	SEG1
ATOM	1192	SG	CYS	78	-8.194	10.098	-7.840	1.00	1.03	SEG1
ATOM	1193	HG	CYS	78	-9.034	9.895	-8.259	1.00	1.50	SEG1
ATOM	1194	C	CYS	78	-8.869	7.094	-8.807	1.00	0.53	SEG1
ATOM	1195	O	CYS	78	-9.898	7.525	-9.293	1.00	0.60	SEG1
ATOM	1196	N	GLY	79	-8.051	6.342	-9.499	1.00	0.50	SEG1
ATOM	1197	HN	GLY	79	-7.232	6.011	-9.087	1.00	0.52	SEG1
ATOM	1198	CA	GLY	79	-8.359	5.990	-10.917	1.00	0.52	SEG1
ATOM	1199	HA1	GLY	79	-7.461	6.072	-11.510	1.00	0.57	SEG1
ATOM	1200	HA2	GLY	79	-9.106	6.671	-11.300	1.00	0.57	SEG1
ATOM	1201	C	GLY	79	-8.887	4.557	-10.998	1.00	0.44	SEG1
ATOM	1202	O	GLY	79	-8.355	3.654	-10.379	1.00	0.39	SEG1
ATOM	1203	N	ARG	80	-9.926	4.344	-11.762	1.00	0.48	SEG1
ATOM	1204	HN	ARG	80	-10.329	5.093	-12.252	1.00	0.54	SEG1
ATOM	1205	CA	ARG	80	-10.498	2.968	-11.900	1.00	0.47	SEG1
ATOM	1206	HA	ARG	80	-10.700	2.549	-10.927	1.00	0.46	SEG1
ATOM	1207	CB	ARG	80	-11.810	3.158	-12.668	1.00	0.57	SEG1
ATOM	1208	HB1	ARG	80	-11.602	3.592	-13.634	1.00	1.05	SEG1
ATOM	1209	HB2	ARG	80	-12.461	3.816	-12.110	1.00	1.00	SEG1
ATOM	1210	CG	ARG	80	-12.497	1.804	-12.859	1.00	1.37	SEG1
ATOM	1211	HG1	ARG	80	-12.708	1.367	-11.894	1.00	1.87	SEG1
ATOM	1212	HG2	ARG	80	-11.846	1.147	-13.417	1.00	1.95	SEG1
ATOM	1213	CD	ARG	80	-13.806	1.999	-13.626	1.00	1.53	SEG1
ATOM	1214	FD1	ARG	80	-13.604	2.294	-14.644	1.00	1.92	SEG1
ATOM	1215	HD2	ARG	80	-14.421	2.740	-13.134	1.00	1.71	SEG1
ATOM	1216	NE	ARG	80	-14.477	0.662	-13.605	1.00	2.30	SEG1
ATOM	1217	HE	ARG	80	-14.158	-0.027	-12.984	1.00	2.85	SEG1
ATOM	1218	CZ	ARG	80	-15.495	0.392	-14.397	1.00	2.84	SEG1
ATOM	1219	NH1	ARG	80	-16.058	-0.783	-14.331	1.00	3.87	SEG1

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ATOM	1210	HH11	APG	80	-15.716	-1.465	-13.685	1.00	4.32	SEG1
ATOM	1211	HH12	APG	80	-16.832	-1.001	-14.926	1.00	4.40	SEG1
ATOM	1212	NH2	APG	80	-15.953	1.276	-15.255	1.00	2.85	SEG1
ATOM	1223	HH21	APG	80	-15.533	2.181	-15.325	1.00	2.56	SEG1
ATOM	1224	HH22	APG	80	-16.729	1.042	-15.843	1.00	3.51	SEG1
ATOM	1225	C	ARG	80	-9.537	2.062	-12.685	1.00	0.43	SEG1
ATOM	1226	O	ARG	80	-9.369	0.900	-12.366	1.00	0.41	SEG1
ATOM	1227	N	GLN	81	-8.919	2.584	-13.715	1.00	0.45	SEG1
ATOM	1228	NN	GLN	81	-9.082	3.521	-13.954	1.00	0.50	SEG1
ATOM	1229	CA	GLN	81	-7.978	1.757	-14.538	1.00	0.45	SEG1
ATOM	1230	HA	GLN	81	-8.518	0.952	-15.012	1.00	0.47	SEG1
ATOM	1231	CB	GLN	81	-7.437	2.708	-15.611	1.00	0.51	SEG1
ATOM	1232	HB1	GLN	81	-6.520	2.310	-16.019	1.00	0.53	SEG1
ATOM	1233	HB2	GLN	81	-7.248	3.676	-15.172	1.00	0.51	SEG1
ATOM	1234	CG	GLN	81	-8.474	2.849	-16.729	1.00	0.57	SEG1
ATOM	1235	HG1	GLN	81	-9.329	3.394	-16.358	1.00	0.93	SEG1
ATOM	1236	HG2	GLN	81	-8.787	1.867	-17.054	1.00	1.00	SEG1
ATOM	1237	CD	GLN	81	-7.863	3.607	-17.909	1.00	1.27	SEG1
ATOM	1238	OE1	GLN	81	-6.983	4.425	-17.732	1.00	1.91	SEG1
ATOM	1239	HE2	GLN	81	-8.299	3.368	-19.116	1.00	2.02	SEG1
ATOM	1240	HE21	GLN	81	-9.010	2.708	-19.259	1.00	2.34	SEG1
ATOM	1241	HE22	GLN	81	-7.917	3.849	-19.880	1.00	2.61	SEG1
ATOM	1242	C	GLN	81	-6.834	1.182	-13.678	1.00	0.41	SEG1
ATOM	1243	O	GLN	81	-6.637	-0.018	-13.664	1.00	0.40	SEG1
ATOM	1244	N	PRO	82	-6.113	2.035	-12.975	1.00	0.40	SEG1
ATOM	1245	CA	PRO	82	-5.002	1.541	-12.119	1.00	0.39	SEG1
ATOM	1246	HA	PRO	82	-4.301	0.972	-12.709	1.00	0.41	SEG1
ATOM	1247	CB	PRO	82	-4.332	2.818	-11.614	1.00	0.42	SEG1
ATOM	1248	HB1	PRO	82	-3.488	3.078	-12.233	1.00	0.47	SEG1
ATOM	1249	HB2	PRO	82	-4.023	2.698	-10.584	1.00	0.42	SEG1
ATOM	1250	CG	PRO	82	-5.382	3.868	-11.719	1.00	0.43	SEG1
ATOM	1251	HG1	PRO	82	-4.921	4.825	-11.895	1.00	0.47	SEG1
ATOM	1252	HG2	PRO	82	-5.973	3.894	-10.814	1.00	0.41	SEG1
ATOM	1253	CD	PRO	82	-6.244	3.499	-12.896	1.00	0.44	SEG1
ATOM	1254	HD2	PRO	82	-7.265	3.790	-12.715	1.00	0.45	SEG1
ATOM	1255	HD1	PRO	82	-5.868	3.956	-13.798	1.00	0.48	SEG1
ATOM	1256	C	PRO	82	-5.539	0.688	-10.964	1.00	0.35	SEG1
ATOM	1257	O	PRO	82	-4.823	-0.122	-10.404	1.00	0.38	SEG1
ATOM	1258	N	CYS	83	-6.793	0.847	-10.614	1.00	0.31	SEG1
ATOM	1259	NN	CYS	83	-7.356	1.496	-11.086	1.00	0.32	SEG1
ATOM	1260	CA	CYS	83	-7.370	0.024	-9.507	1.00	0.30	SEG1
ATOM	1261	HA	CYS	83	-6.782	0.134	-8.608	1.00	0.31	SEG1
ATOM	1262	CB	CYS	83	-8.783	0.568	-9.282	1.00	0.34	SEG1
ATOM	1263	HB1	CYS	83	-9.409	0.301	-10.121	1.00	0.66	SEG1
ATOM	1264	HB2	CYS	83	-8.745	1.643	-9.188	1.00	0.77	SEG1
ATOM	1265	SG	CYS	83	-9.469	-0.147	-7.767	1.00	0.99	SEG1
ATOM	1266	HG	CYS	83	-10.419	-0.228	-7.881	1.00	1.67	SEG1
ATOM	1267	C	CYS	83	-7.422	-1.444	-9.938	1.00	0.29	SEG1
ATOM	1268	O	CYS	83	-7.020	-2.331	-9.209	1.00	0.28	SEG1
ATOM	1269	N	GLY	84	-7.908	-1.700	-11.129	1.00	0.30	SEG1
ATOM	1270	NN	GLY	84	-8.216	-0.962	-11.696	1.00	0.32	SEG1
ATOM	1271	CA	GLY	84	-7.984	-3.104	-11.635	1.00	0.31	SEG1
ATOM	1272	HA1	GLY	84	-8.409	-3.109	-12.627	1.00	0.33	SEG1
ATOM	1273	HA2	GLY	84	-8.606	-3.691	-10.973	1.00	0.31	SEG1
ATOM	1274	C	GLY	84	-6.579	-3.706	-11.684	1.00	0.30	SEG1
ATOM	1275	O	GLY	84	-6.374	-4.861	-11.358	1.00	0.30	SEG1
ATOM	1276	N	ARG	85	-5.609	-2.928	-12.090	1.00	0.31	SEG1
ATOM	1277	NN	ARG	85	-5.804	-2.002	-12.347	1.00	0.32	SEG1
ATOM	1278	CA	ARG	85	-4.209	-3.444	-12.158	1.00	0.32	SEG1
ATOM	1279	HA	ARG	85	-4.174	-4.358	-12.731	1.00	0.34	SEG1
ATOM	1280	CB	ARG	85	-3.404	-2.351	-12.866	1.00	0.35	SEG1
ATOM	1281	HB1	ARG	85	-2.351	-2.578	-12.797	1.00	0.70	SEG1
ATOM	1282	HB2	ARG	85	-3.600	-1.398	-12.396	1.00	0.87	SEG1
ATOM	1283	CG	ARG	85	-3.814	-2.286	-14.339	1.00	0.87	SEG1
ATOM	1284	HG1	ARG	85	-4.866	-2.058	-14.411	1.00	1.41	SEG1
ATOM	1285	HG2	ARG	85	-3.618	-3.239	-14.810	1.00	1.40	SEG1
ATOM	1286	CD	ARG	85	-3.009	-1.193	-15.048	1.00	1.12	SEG1
ATOM	1287	HD1	ARG	85	-1.958	-1.439	-15.049	1.00	1.59	SEG1
ATOM	1288	HD2	ARG	85	-3.174	-0.237	-14.569	1.00	1.64	SEG1
ATOM	1289	NE	ARG	85	-3.530	-1.173	-16.451	1.00	1.63	SEG1
ATOM	1290	HE	ARG	85	-4.064	-1.928	-16.777	1.00	2.16	SEG1
ATOM	1291	CZ	ARG	85	-3.292	-0.158	-17.259	1.00	2.06	SEG1
ATOM	1292	NH1	ARG	85	-3.780	-0.179	-18.469	1.00	2.88	SEG1
ATOM	1293	HH11	ARG	85	-4.327	-0.959	-18.773	1.00	3.33	SEG1
ATOM	1294	HH12	ARG	85	-3.608	0.584	-19.091	1.00	3.28	SEG1
ATOM	1295	NH2	ARG	85	-2.572	0.373	-16.877	1.00	2.26	SEG1
ATOM	1296	HH21	ARG	85	-2.186	0.908	-15.956	1.00	2.26	SEG1

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ATOM	1307	HN2	ARG	85	-2.411	-5.628	-17.513	1.00	2.81	SEG1
ATOM	1308	C	ARG	85	-3.672	-5.680	-10.741	1.00	0.29	SEG1
ATOM	1309	O	ARG	85	-2.915	-4.600	-10.501	1.00	0.29	SEG1
ATOM	1310	N	PHE	86	-4.055	-1.643	-9.808	1.00	0.27	SEG1
ATOM	1311	NN	PHE	86	-4.661	-2.106	-10.038	1.00	0.28	SEG1
ATOM	1312	CA	PHE	86	-3.563	-1.999	-8.460	1.00	0.26	SEG1
ATOM	1313	HA	PHE	86	-2.493	-2.880	-8.380	1.00	0.28	SEG1
ATOM	1314	CB	PHE	86	-4.245	-1.859	-7.626	1.00	0.27	SEG1
ATOM	1315	HB1	PHE	86	-5.317	-1.950	-7.727	1.00	0.28	SEG1
ATOM	1316	HB2	PHE	86	-3.927	-0.911	-8.034	1.00	0.28	SEG1
ATOM	1317	CG	PHE	86	-3.875	-1.918	-6.159	1.00	0.25	SEG1
ATOM	1318	CD1	PHE	86	-4.611	-2.735	-5.292	1.00	1.23	SEG1
ATOM	1319	HD1	PHE	86	-5.430	-3.327	-5.675	1.00	2.16	SEG1
ATOM	1311	CD2	PHE	86	-2.813	-1.149	-5.663	1.00	1.24	SEG1
ATOM	1311	HD2	PHE	86	-2.242	-0.517	-6.330	1.00	2.17	SEG1
ATOM	1311	CE1	PHE	86	-4.286	-2.768	-3.932	1.00	1.23	SEG1
ATOM	1311	HE1	PHE	86	-4.855	-3.420	-3.255	1.00	2.16	SEG1
ATOM	1314	CE2	PHE	86	-2.488	-1.203	-4.300	1.00	1.24	SEG1
ATOM	1315	HE2	PHE	86	-1.671	-0.609	-3.914	1.00	2.17	SEG1
ATOM	1316	CZ	PHE	86	-3.225	-2.022	-3.435	1.00	0.26	SEG1
ATOM	1317	HZ	PHE	86	-2.974	-2.063	-2.385	1.00	0.28	SEG1
ATOM	1318	C	PHE	86	-3.973	-4.364	-7.622	1.00	0.25	SEG1
ATOM	1319	O	PHE	86	-3.152	-5.079	-7.278	1.00	0.27	SEG1
ATOM	1320	N	LEU	87	-5.229	-4.726	-7.926	1.00	0.25	SEG1
ATOM	1321	NN	LEU	87	-5.875	-4.131	-8.358	1.00	0.25	SEG1
ATOM	1322	CA	LEU	87	-5.673	-6.045	-7.366	1.00	0.25	SEG1
ATOM	1323	HA	LEU	87	-5.279	-6.160	-6.369	1.00	0.26	SEG1
ATOM	1324	CB	LEU	87	-7.210	-5.988	-7.297	1.00	0.26	SEG1
ATOM	1325	HB1	LEU	87	-7.506	-5.182	-6.641	1.00	0.27	SEG1
ATOM	1326	HB2	LEU	87	-7.580	-6.922	-6.899	1.00	0.28	SEG1
ATOM	1327	CG	LEU	87	-7.818	-5.758	-8.685	1.00	0.28	SEG1
ATOM	1328	HG	LEU	87	-7.108	-5.253	-9.314	1.00	0.28	SEG1
ATOM	1329	CD1	LEU	87	-8.194	-7.100	-9.314	1.00	0.32	SEG1
ATOM	1330	HD11	LEU	87	-9.169	-7.402	-8.963	1.00	1.08	SEG1
ATOM	1331	HD12	LEU	87	-7.465	-7.846	-9.036	1.00	1.03	SEG1
ATOM	1332	HD13	LEU	87	-8.215	-7.000	-10.389	1.00	1.05	SEG1
ATOM	1333	CD2	LEU	87	-9.078	-4.897	-8.551	1.00	0.30	SEG1
ATOM	1334	HD21	LEU	87	-8.798	-3.887	-8.293	1.00	1.03	SEG1
ATOM	1335	HD22	LEU	87	-9.711	-5.304	-7.778	1.00	1.02	SEG1
ATOM	1336	HD23	LEU	87	-9.612	-4.893	-9.490	1.00	1.08	SEG1
ATOM	1337	C	LEU	87	-5.191	-7.214	-8.241	1.00	0.25	SEG1
ATOM	1338	O	LEU	87	-4.769	-8.240	-7.732	1.00	0.27	SEG1
ATOM	1339	N	ARG	88	-5.240	-7.072	-9.544	1.00	0.26	SEG1
ATOM	1340	NN	ARG	88	-5.573	-6.239	-9.934	1.00	0.26	SEG1
ATOM	1341	CA	ARG	88	-4.777	-8.183	-10.434	1.00	0.27	SEG1
ATOM	1342	HA	ARG	88	-5.320	-9.088	-10.219	1.00	0.28	SEG1
ATOM	1343	CB	ARG	88	-5.079	-7.716	-11.860	1.00	0.29	SEG1
ATOM	1344	HB1	ARG	88	-4.464	-6.862	-12.099	1.00	0.79	SEG1
ATOM	1345	HB2	ARG	88	-6.122	-7.443	-11.936	1.00	0.79	SEG1
ATOM	1346	CG	ARG	88	-4.774	-8.850	-12.841	1.00	1.08	SEG1
ATOM	1347	HG1	ARG	88	-5.390	-9.705	-12.604	1.00	1.57	SEG1
ATOM	1348	HG2	ARG	88	-3.732	-9.123	-12.761	1.00	1.56	SEG1
ATOM	1349	CD	ARG	88	-5.075	-8.391	-14.270	1.00	1.09	SEG1
ATOM	1350	HD1	ARG	88	-4.345	-7.665	-14.593	1.00	1.19	SEG1
ATOM	1351	HD2	ARG	88	-6.072	-7.977	-14.328	1.00	1.44	SEG1
ATOM	1352	NE	ARG	88	-4.972	-9.628	-15.109	1.00	2.05	SEG1
ATOM	1353	HE	ARG	88	-4.837	-10.496	-14.674	1.00	2.70	SEG1
ATOM	1354	CZ	ARG	88	-5.058	-9.575	-16.423	1.00	2.54	SEG1
ATOM	1355	NH1	ARG	88	-4.962	-10.680	-17.111	1.00	3.60	SEG1
ATOM	1356	NH11	ARG	88	-4.825	-11.552	-16.640	1.00	4.10	SEG1
ATOM	1357	NH12	ARG	88	-5.026	-10.655	-18.108	1.00	4.07	SEG1
ATOM	1358	NH2	ARG	88	-5.240	-8.439	-17.056	1.00	2.48	SEG1
ATOM	1359	NH21	ARG	88	-5.318	-7.583	-16.548	1.00	2.17	SEG1
ATOM	1360	NH22	ARG	88	-5.301	-8.432	-18.054	1.00	3.15	SEG1
ATOM	1361	C	ARG	88	-3.278	-8.413	-10.244	1.00	0.28	SEG1
ATOM	1362	O	ARG	88	-2.823	-9.536	-10.140	1.00	0.30	SEG1
ATOM	1363	N	ALA	89	-2.514	-7.354	-10.182	1.00	0.29	SEG1
ATOM	1364	NN	ALA	89	-2.912	-6.461	-10.258	1.00	0.29	SEG1
ATOM	1365	CA	ALA	89	-1.041	-7.500	-9.980	1.00	0.31	SEG1
ATOM	1366	HA	ALA	89	-0.633	-8.202	-10.690	1.00	0.33	SEG1
ATOM	1367	CB	ALA	89	-0.457	-6.110	-10.226	1.00	0.35	SEG1
ATOM	1368	HB1	ALA	89	0.588	-6.201	-10.431	1.00	1.03	SEG1
ATOM	1369	HB2	ALA	89	-0.560	-5.513	-9.331	1.00	1.04	SEG1
ATOM	1370	HB3	ALA	89	-0.986	-5.635	-11.039	1.00	1.09	SEG1
ATOM	1371	C	ALA	89	-0.750	-7.957	-9.549	1.00	0.30	SEG1
ATOM	1372	O	ALA	89	0.284	-8.537	-8.273	1.00	0.31	SEG1
ATOM	1373	N	TYR	90	-1.651	-7.689	-7.634	1.00	0.30	SEG1

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ATOM	1374	HN	TYR	90	-2.471	-7.213	-7.881	1.00	0.30	SEG1
ATOM	1375	CA	TYR	90	-1.427	-8.096	-6.114	1.00	0.31	SEG1
ATOM	1376	HA	TYR	90	-0.554	-7.599	-5.819	1.00	0.34	SEG1
ATOM	1377	CB	TYR	90	-2.676	-7.634	-5.658	1.00	0.33	SEG1
ATOM	1378	HB1	TYR	90	-3.544	-8.133	-5.859	1.00	0.41	SEG1
ATOM	1379	HB2	TYR	90	-2.790	-6.565	-5.572	1.00	0.34	SEG1
ATOM	1380	CG	TYR	90	-2.535	-7.972	-3.994	1.00	0.44	SEG1
ATOM	1381	CD1	TYR	90	-1.951	-7.050	-3.117	1.00	1.36	SEG1
ATOM	1382	HD1	TYR	90	-1.603	-6.097	-3.488	1.00	2.21	SEG1
ATOM	1383	CD2	TYR	90	-2.984	-9.207	-3.514	1.00	1.28	SEG1
ATOM	1384	HD2	TYR	90	-3.436	-9.918	-4.191	1.00	2.18	SEG1
ATOM	1385	CE1	TYR	90	-1.818	-7.363	-1.761	1.00	1.49	SEG1
ATOM	1386	HE1	TYR	90	-1.367	-6.653	-1.084	1.00	2.40	SEG1
ATOM	1387	CE2	TYR	90	-2.850	-9.520	-2.158	1.00	1.32	SEG1
ATOM	1388	HE2	TYR	90	-3.192	-10.475	-1.788	1.00	2.17	SEG1
ATOM	1389	CZ	TYR	90	-2.266	-8.597	-1.282	1.00	0.83	SEG1
ATOM	1390	OH	TYR	90	-2.139	-8.902	0.055	1.00	1.04	SEG1
ATOM	1391	HH	TYR	90	-1.881	-9.824	0.126	1.00	1.37	SEG1
ATOM	1392	C	TYR	90	-1.263	-9.617	-6.114	1.00	0.31	SEG1
ATOM	1393	O	TYR	90	-0.349	-10.104	-5.475	1.00	0.33	SEG1
ATOM	1394	N	ARG	91	-2.138	-10.373	-6.741	1.00	0.31	SEG1
ATOM	1395	HN	ARG	91	-2.868	-9.958	-7.252	1.00	0.32	SEG1
ATOM	1396	CA	ARG	91	-2.015	-11.866	-6.674	1.00	0.34	SEG1
ATOM	1397	HA	ARG	91	-2.099	-12.206	-5.657	1.00	0.38	SEG1
ATOM	1398	CB	ARG	91	-3.175	-12.418	-7.509	1.00	0.39	SEG1
ATOM	1399	HB1	ARG	91	-3.089	-12.066	-8.525	1.00	0.91	SEG1
ATOM	1400	HB2	ARG	91	-4.111	-12.081	-7.087	1.00	0.81	SEG1
ATOM	1401	CG	ARG	91	-3.130	-13.950	-7.493	1.00	1.12	SEG1
ATOM	1402	HG1	ARG	91	-3.219	-14.301	-6.475	1.00	1.64	SEG1
ATOM	1403	HG2	ARG	91	-2.190	-14.285	-7.908	1.00	1.67	SEG1
ATOM	1404	CD	ARG	91	-4.287	-14.512	-8.325	1.00	1.16	SEG1
ATOM	1405	HD1	ARG	91	-4.191	-14.204	-9.355	1.00	1.49	SEG1
ATOM	1406	HD2	ARG	91	-5.235	-14.185	-7.920	1.00	1.29	SEG1
ATOM	1407	NE	ARG	91	-4.160	-16.002	-8.224	1.00	2.06	SEG1
ATOM	1408	HE	ARG	91	-3.413	-16.391	-7.720	1.00	2.61	SEG1
ATOM	1409	CZ	ARG	91	-5.036	-16.806	-8.796	1.00	2.58	SEG1
ATOM	1410	NH1	ARG	91	-4.891	-18.097	-8.674	1.00	3.56	SEG1
ATOM	1411	HH11	ARG	91	-4.124	-18.466	-8.151	1.00	4.00	SEG1
ATOM	1412	HH12	ARG	91	-5.549	-18.716	-9.103	1.00	4.01	SEG1
ATOM	1413	NH2	ARG	91	-6.050	-16.338	-9.487	1.00	2.55	SEG1
ATOM	1414	HH21	ARG	91	-6.180	-15.354	-9.595	1.00	2.26	SEG1
ATOM	1415	HH22	ARG	91	-6.698	-16.973	-9.908	1.00	3.16	SEG1
ATOM	1416	C	ARG	91	-0.668	-12.297	-7.258	1.00	0.32	SEG1
ATOM	1417	O	ARG	91	-0.004	-13.170	-6.730	1.00	0.33	SEG1
ATOM	1418	N	GLU	92	-0.251	-11.671	-8.328	1.00	0.30	SEG1
ATOM	1419	HN	GLU	92	-0.802	-10.957	-8.721	1.00	0.31	SEG1
ATOM	1420	CA	GLU	92	1.070	-12.021	-8.933	1.00	0.32	SEG1
ATOM	1421	HA	GLU	92	1.090	-13.061	-9.213	1.00	0.34	SEG1
ATOM	1422	CB	GLU	92	1.201	-11.139	-10.177	1.00	0.38	SEG1
ATOM	1423	HB1	GLU	92	2.205	-11.212	-10.567	1.00	0.42	SEG1
ATOM	1424	HB2	GLU	92	0.989	-10.112	-9.914	1.00	0.38	SEG1
ATOM	1425	CG	GLU	92	0.206	-11.612	-11.243	1.00	0.42	SEG1
ATOM	1426	HG1	GLU	92	-0.798	-11.545	-10.851	1.00	0.56	SEG1
ATOM	1427	HG2	GLU	92	0.421	-12.638	-11.500	1.00	0.63	SEG1
ATOM	1428	CD	GLU	92	0.317	-10.739	-12.503	1.00	0.87	SEG1
ATOM	1429	OE1	GLU	92	-0.277	-11.108	-13.503	1.00	1.48	SEG1
ATOM	1430	OE2	GLU	92	0.983	-9.715	-12.449	1.00	1.58	SEG1
ATOM	1431	C	GLU	92	2.187	-11.716	-7.931	1.00	0.33	SEG1
ATOM	1432	O	GLU	92	3.138	-12.462	-7.796	1.00	0.35	SEG1
ATOM	1433	N	GLY	93	2.068	-10.618	-7.229	1.00	0.35	SEG1
ATOM	1434	HN	GLY	93	1.289	-10.040	-7.363	1.00	0.35	SEG1
ATOM	1435	CA	GLY	93	3.107	-10.238	-6.226	1.00	0.40	SEG1
ATOM	1436	HA1	GLY	93	2.839	-9.297	-5.770	1.00	0.45	SEG1
ATOM	1437	HA2	GLY	93	4.062	-10.137	-6.722	1.00	0.43	SEG1
ATOM	1438	C	GLY	93	3.208	-11.314	-5.141	1.00	0.37	SEG1
ATOM	1439	O	GLY	93	4.291	-11.691	-4.733	1.00	0.39	SEG1
ATOM	1440	N	ALA	94	2.089	-11.816	-4.671	1.00	0.37	SEG1
ATOM	1441	HN	ALA	94	1.229	-11.498	-5.017	1.00	0.37	SEG1
ATOM	1442	CA	ALA	94	2.125	-12.877	-3.611	1.00	0.41	SEG1
ATOM	1443	HA	ALA	94	2.591	-12.499	-2.714	1.00	0.46	SEG1
ATOM	1444	CB	ALA	94	0.661	-13.227	-3.336	1.00	0.47	SEG1
ATOM	1445	HB1	ALA	94	0.613	-14.032	-2.617	1.00	1.13	SEG1
ATOM	1446	HB2	ALA	94	0.186	-13.536	-4.255	1.00	1.16	SEG1
ATOM	1447	HB3	ALA	94	0.152	-12.361	-2.941	1.00	1.08	SEG1
ATOM	1448	C	ALA	94	2.882	-14.100	-4.134	1.00	0.36	SEG1
ATOM	1449	O	ALA	94	3.673	-14.708	-3.435	1.00	0.39	SEG1
ATOM	1450	N	LEU	95	2.654	-14.446	-5.373	1.00	0.32	SEG1

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ATOM	1451	HN	LEU	95	2.023	-13.923	-5.914	1.00	0.31	SEG1
ATOM	1452	CA	LEU	95	3.364	-15.612	-5.978	1.00	0.34	SEG1
ATOM	1453	HA	LEU	95	3.152	-16.510	-5.423	1.00	0.39	SEG1
ATOM	1454	CB	LEU	95	2.804	-15.722	-7.395	1.00	0.35	SEG1
ATOM	1455	HB1	LEU	95	3.329	-16.496	-7.933	1.00	0.40	SEG1
ATOM	1456	HB2	LEU	95	2.926	-14.777	-7.907	1.00	0.33	SEG1
ATOM	1457	CG	LEU	95	1.318	-16.077	-7.312	1.00	0.39	SEG1
ATOM	1458	HG	LEU	95	0.885	-15.596	-6.446	1.00	0.39	SEG1
ATOM	1459	CD1	LEU	95	0.591	-15.596	-8.574	1.00	0.39	SEG1
ATOM	1460	HD11	LEU	95	-0.039	-16.386	-8.955	1.00	1.13	SEG1
ATOM	1461	HD12	LEU	95	1.314	-15.318	-9.326	1.00	1.12	SEG1
ATOM	1462	HD13	LEU	95	-0.020	-14.738	-8.328	1.00	1.04	SEG1
ATOM	1463	CD2	LEU	95	1.170	-17.596	-7.176	1.00	0.55	SEG1
ATOM	1464	HD21	LEU	95	1.790	-18.086	-7.912	1.00	1.27	SEG1
ATOM	1465	HD22	LEU	95	0.137	-17.873	-7.332	1.00	1.12	SEG1
ATOM	1466	HD23	LEU	95	1.478	-17.901	-6.186	1.00	1.12	SEG1
ATOM	1467	C	LEU	95	4.867	-15.330	-5.990	1.00	0.33	SEG1
ATOM	1468	O	LEU	95	5.677	-16.193	-5.713	1.00	0.38	SEG1
ATOM	1469	N	ARG	96	5.235	-14.111	-6.293	1.00	0.32	SEG1
ATOM	1470	HN	ARG	96	4.553	-13.438	-6.497	1.00	0.31	SEG1
ATOM	1471	CA	ARG	96	6.681	-13.734	-6.311	1.00	0.39	SEG1
ATOM	1472	HA	ARG	96	7.212	-14.310	-7.051	1.00	0.43	SEG1
ATOM	1473	CB	ARG	96	6.715	-12.247	-6.679	1.00	0.46	SEG1
ATOM	1474	HB1	ARG	96	6.207	-11.676	-5.917	1.00	1.01	SEG1
ATOM	1475	HB2	ARG	96	6.223	-12.099	-7.629	1.00	0.84	SEG1
ATOM	1476	CG	ARG	96	8.170	-11.783	-6.775	1.00	1.29	SEG1
ATOM	1477	HG1	ARG	96	8.680	-12.356	-7.535	1.00	1.76	SEG1
ATOM	1478	HG2	ARG	96	8.657	-11.934	-5.824	1.00	2.06	SEG1
ATOM	1479	CD	ARG	96	8.216	-10.299	-7.143	1.00	1.52	SEG1
ATOM	1480	HD1	ARG	96	9.233	-9.938	-7.124	1.00	1.96	SEG1
ATOM	1481	HD2	ARG	96	7.599	-9.726	-6.465	1.00	1.67	SEG1
ATOM	1482	NE	ARG	96	7.673	-10.230	-8.534	1.00	2.44	SEG1
ATOM	1483	HE	ARG	96	6.756	-9.913	-8.674	1.00	2.95	SEG1
ATOM	1484	CZ	ARG	96	8.400	-10.582	-9.576	1.00	3.12	SEG1
ATOM	1485	NH1	ARG	96	7.881	-10.497	-10.771	1.00	4.14	SEG1
ATOM	1486	NH11	ARG	96	6.944	-10.169	-10.887	1.00	4.47	SEG1
ATOM	1487	NH12	ARG	96	8.421	-10.762	-11.571	1.00	4.77	SEG1
ATOM	1488	NH2	ARG	96	9.632	-11.019	-9.440	1.00	3.27	SEG1
ATOM	1489	NH21	ARG	96	10.046	-11.094	-8.534	1.00	2.90	SEG1
ATOM	1490	NH22	ARG	96	10.159	-11.279	-10.249	1.00	4.07	SEG1
ATOM	1491	C	ARG	96	7.289	-13.955	-4.922	1.00	0.40	SEG1
ATOM	1492	O	ARG	96	8.434	-14.341	-4.790	1.00	0.46	SEG1
ATOM	1493	N	ALA	97	6.523	-13.714	-3.884	1.00	0.40	SEG1
ATOM	1494	HN	ALA	97	5.603	-13.403	-4.021	1.00	0.38	SEG1
ATOM	1495	CA	ALA	97	7.048	-13.910	-2.495	1.00	0.46	SEG1
ATOM	1496	HA	ALA	97	7.867	-13.237	-2.302	1.00	0.51	SEG1
ATOM	1497	CB	ALA	97	5.874	-13.592	-1.568	1.00	0.52	SEG1
ATOM	1498	HB1	ALA	97	6.194	-13.678	-0.541	1.00	1.09	SEG1
ATOM	1499	HB2	ALA	97	5.071	-14.289	-1.752	1.00	1.19	SEG1
ATOM	1500	HB3	ALA	97	5.530	-12.586	-1.755	1.00	1.12	SEG1
ATOM	1501	C	ALA	97	7.491	-15.361	-2.312	1.00	0.46	SEG1
ATOM	1502	O	ALA	97	8.513	-15.635	-1.711	1.00	0.50	SEG1
ATOM	1503	N	ALA	98	6.739	-16.288	-2.845	1.00	0.46	SEG1
ATOM	1504	HN	ALA	98	5.928	-16.034	-3.337	1.00	0.43	SEG1
ATOM	1505	CA	ALA	98	7.128	-17.728	-2.728	1.00	0.51	SEG1
ATOM	1506	HA	ALA	98	7.177	-18.023	-1.692	1.00	0.56	SEG1
ATOM	1507	CB	ALA	98	6.029	-18.510	-3.452	1.00	0.55	SEG1
ATOM	1508	HB1	ALA	98	6.167	-18.422	-4.520	1.00	1.20	SEG1
ATOM	1509	HB2	ALA	98	5.064	-18.110	-3.180	1.00	1.10	SEG1
ATOM	1510	HB3	ALA	98	6.081	-19.551	-3.167	1.00	1.06	SEG1
ATOM	1511	C	ALA	98	8.484	-17.938	-3.411	1.00	0.52	SEG1
ATOM	1512	O	ALA	98	9.337	-18.652	-2.922	1.00	0.57	SEG1
ATOM	1513	N	LEU	99	8.684	-17.296	-4.536	1.00	0.50	SEG1
ATOM	1514	HN	LEU	99	7.978	-16.717	-4.895	1.00	0.47	SEG1
ATOM	1515	CA	LEU	99	9.983	-17.421	-5.265	1.00	0.55	SEG1
ATOM	1516	HA	LEU	99	10.159	-18.448	-5.546	1.00	0.60	SEG1
ATOM	1517	CB	LEU	99	9.840	-16.549	-6.517	1.00	0.57	SEG1
ATOM	1518	HB1	LEU	99	9.677	-15.524	-6.223	1.00	0.54	SEG1
ATOM	1519	HB2	LEU	99	9.001	-16.895	-7.101	1.00	0.58	SEG1
ATOM	1520	CG	LEU	99	11.118	-16.638	-7.356	1.00	0.66	SEG1
ATOM	1521	HG	LEU	99	11.953	-16.885	-6.717	1.00	0.67	SEG1
ATOM	1522	CD1	LEU	99	10.956	-17.726	-8.419	1.00	0.73	SEG1
ATOM	1523	HD11	LEU	99	11.553	-17.476	-9.284	1.00	1.16	SEG1
ATOM	1524	HD12	LEU	99	9.917	-17.796	-8.708	1.00	1.14	SEG1
ATOM	1525	HD13	LEU	99	11.282	-18.673	-8.017	1.00	1.42	SEG1
ATOM	1526	CD2	LEU	99	11.376	-15.291	-8.036	1.00	0.70	SEG1
ATOM	1527	HD21	LEU	99	12.164	-15.401	-8.767	1.00	1.16	SEG1

ATOM	1528	HD23	LEU	99	11.673	-14.565	-7.293	1.00	1.32	SEG1
ATOM	1529	HD23	LEU	99	10.474	-14.956	-8.527	1.00	1.12	SEG1
ATOM	1530	C	LEU	99	11.121	-16.906	-4.378	1.00	0.54	SEG1
ATOM	1531	O	LEU	99	12.204	-17.457	-4.362	1.00	0.59	SEG1
ATOM	1532	N	GLN	100	10.881	-15.843	-3.648	1.00	0.50	SEG1
ATOM	1533	HN	GLN	100	9.999	-15.415	-3.636	1.00	0.47	SEG1
ATOM	1534	CA	GLN	100	11.949	-15.274	-2.767	1.00	0.52	SEG1
ATOM	1535	HA	GLN	100	12.788	-14.937	-3.355	1.00	0.57	SEG1
ATOM	1536	CB	GLN	100	11.300	-14.084	-2.049	1.00	0.51	SEG1
ATOM	1537	HB1	GLN	100	11.958	-13.734	-1.269	1.00	0.55	SEG1
ATOM	1538	HB2	GLN	100	10.363	-14.397	-1.614	1.00	0.47	SEG1
ATOM	1539	CG	GLN	100	11.043	-12.947	-3.042	1.00	0.54	SEG1
ATOM	1540	HG1	GLN	100	10.565	-12.126	-2.529	1.00	1.00	SEG1
ATOM	1541	HG2	GLN	100	10.400	-13.301	-3.835	1.00	1.14	SEG1
ATOM	1542	CD	GLN	100	12.369	-12.468	-3.637	1.00	1.30	SEG1
ATOM	1543	DE1	GLN	100	13.079	-11.692	-3.030	1.00	1.83	SEG1
ATOM	1544	NE2	GLN	100	12.739	-12.919	-4.803	1.00	2.26	SEG1
ATOM	1545	HE21	GLN	100	12.169	-13.557	-5.282	1.00	2.51	SEG1
ATOM	1546	HE22	GLN	100	13.583	-12.620	-5.199	1.00	2.99	SEG1
ATOM	1547	C	GLN	100	12.393	-16.320	-1.747	1.00	0.53	SEG1
ATOM	1548	O	GLN	100	13.567	-16.453	-1.455	1.00	0.57	SEG1
ATOM	1549	N	ARG	101	11.464	-17.068	-1.213	1.00	0.51	SEG1
ATOM	1550	HN	ARG	101	10.526	-16.941	-1.475	1.00	0.50	SEG1
ATOM	1551	CA	ARG	101	11.832	-18.118	-0.213	1.00	0.55	SEG1
ATOM	1552	HA	ARG	101	12.287	-17.668	0.656	1.00	0.54	SEG1
ATOM	1553	CB	ARG	101	10.511	-18.789	0.173	1.00	0.58	SEG1
ATOM	1554	HB1	ARG	101	10.061	-19.231	-0.702	1.00	0.62	SEG1
ATOM	1555	HB2	ARG	101	9.841	-18.048	0.587	1.00	0.56	SEG1
ATOM	1556	CG	ARG	101	10.772	-19.878	1.216	1.00	0.64	SEG1
ATOM	1557	HG1	ARG	101	11.219	-19.436	2.094	1.00	0.62	SEG1
ATOM	1558	HG2	ARG	101	11.444	-20.617	0.803	1.00	0.67	SEG1
ATOM	1559	CD	ARG	101	9.450	-20.545	1.601	1.00	0.70	SEG1
ATOM	1560	HD1	ARG	101	9.077	-21.143	0.784	1.00	1.17	SEG1
ATOM	1561	HD2	ARG	101	8.724	-19.797	1.885	1.00	1.21	SEG1
ATOM	1562	NE	ARG	101	9.785	-21.423	2.766	1.00	1.42	SEG1
ATOM	1563	HE	ARG	101	10.714	-21.706	2.902	1.00	1.98	SEG1
ATOM	1564	CZ	ARG	101	8.861	-21.815	3.621	1.00	1.81	SEG1
ATOM	1565	NH1	ARG	101	9.209	-22.561	4.633	1.00	2.84	SEG1
ATOM	1566	HH11	ARG	101	10.165	-22.827	4.753	1.00	3.38	SEG1
ATOM	1567	HH12	ARG	101	8.519	-22.866	5.290	1.00	3.17	SEG1
ATOM	1568	NH2	ARG	101	7.599	-21.479	3.475	1.00	1.50	SEG1
ATOM	1569	HH21	ARG	101	7.310	-20.915	2.704	1.00	1.32	SEG1
ATOM	1570	HH22	ARG	101	6.923	-21.791	4.143	1.00	1.89	SEG1
ATOM	1571	C	ARG	101	12.794	-19.130	-0.855	1.00	0.63	SEG1
ATOM	1572	O	ARG	101	13.839	-19.433	-0.312	1.00	0.66	SEG1
ATOM	1573	N	SER	102	12.450	-19.639	-2.014	1.00	0.67	SEG1
ATOM	1574	HN	SER	102	11.606	-19.368	-2.433	1.00	0.65	SEG1
ATOM	1575	CA	SER	102	13.350	-20.619	-2.703	1.00	0.76	SEG1
ATOM	1576	HA	SER	102	13.558	-21.458	-2.058	1.00	0.81	SEG1
ATOM	1577	CB	SER	102	12.571	-21.083	-3.935	1.00	0.80	SEG1
ATOM	1578	HB1	SER	102	13.223	-21.667	-4.571	1.00	1.16	SEG1
ATOM	1579	HB2	SER	102	12.213	-20.227	-4.482	1.00	1.31	SEG1
ATOM	1580	OG	SER	102	11.462	-21.870	-3.520	1.00	1.48	SEG1
ATOM	1581	HG	SER	102	10.933	-21.343	-2.918	1.00	1.81	SEG1
ATOM	1582	C	SER	102	14.652	-19.929	-3.125	1.00	0.78	SEG1
ATOM	1583	O	SER	102	15.727	-20.489	-3.014	1.00	0.85	SEG1
ATOM	1584	N	LEU	103	14.556	-18.713	-3.599	1.00	0.74	SEG1
ATOM	1585	HN	LEU	103	13.676	-18.287	-3.667	1.00	0.69	SEG1
ATOM	1586	CA	LEU	103	15.779	-17.963	-4.027	1.00	0.79	SEG1
ATOM	1587	HA	LEU	103	16.362	-18.562	-4.710	1.00	0.86	SEG1
ATOM	1588	CB	LEU	103	15.262	-16.712	-4.742	1.00	0.77	SEG1
ATOM	1589	HB1	LEU	103	15.875	-15.865	-4.473	1.00	0.80	SEG1
ATOM	1590	HB2	LEU	103	14.240	-16.525	-4.448	1.00	0.71	SEG1
ATOM	1591	CG	LEU	103	15.328	-16.923	-6.257	1.00	0.83	SEG1
ATOM	1592	HG	LEU	103	16.340	-17.176	-6.532	1.00	0.90	SEG1
ATOM	1593	CD1	LEU	103	14.391	-18.063	-6.668	1.00	0.82	SEG1
ATOM	1594	HD11	LEU	103	13.618	-18.179	-5.926	1.00	1.34	SEG1
ATOM	1595	HD12	LEU	103	14.955	-18.980	-6.746	1.00	1.29	SEG1
ATOM	1596	HD13	LEU	103	13.943	-17.836	-7.623	1.00	1.23	SEG1
ATOM	1597	CD2	LEU	103	14.909	-15.636	-6.970	1.00	0.84	SEG1
ATOM	1598	HD21	LEU	103	14.934	-15.792	-8.038	1.00	1.29	SEG1
ATOM	1599	HD22	LEU	103	15.591	-14.841	-6.706	1.00	1.31	SEG1
ATOM	1600	HD23	LEU	103	13.907	-15.366	-6.668	1.00	1.28	SEG1
ATOM	1601	C	LEU	103	16.623	-17.580	-2.813	1.00	0.79	SEG1
ATOM	1602	O	LEU	103	17.820	-17.386	-2.917	1.00	0.87	SEG1
ATOM	1603	N	ALA	104	16.009	-17.465	-1.662	1.00	0.72	SEG1
ATOM	1604	HN	ALA	104	15.044	-17.623	-1.611	1.00	0.67	SEG1

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ATOM	1605	CA	ALA	104	16.772	-17.089	-0.408	1.00	0.74	SEG1
ATOM	1606	HA	ALA	104	17.126	-16.074	-0.497	1.00	0.76	SEG1
ATOM	1607	CB	ALA	104	15.765	-17.216	0.717	1.00	0.68	SEG1
ATOM	1608	HB1	ALA	104	15.952	-16.444	1.449	1.00	1.17	SEG1
ATOM	1609	HB2	ALA	104	15.869	-18.185	1.142	1.00	1.16	SEG1
ATOM	1610	HB3	ALA	104	14.763	-17.109	0.328	1.00	1.31	SEG1
ATOM	1611	C	ALA	104	17.944	-18.055	-0.218	1.00	0.83	SEG1
ATOM	1612	O	ALA	104	19.073	-17.641	-0.042	1.00	0.90	SEG1
ATOM	1613	N	ALA	105	17.681	-19.335	-0.254	1.00	0.85	SEG1
ATOM	1614	HN	ALA	105	16.762	-19.639	-0.411	1.00	0.81	SEG1
ATOM	1615	CA	ALA	105	18.779	-20.333	-0.073	1.00	0.95	SEG1
ATOM	1616	HA	ALA	105	19.323	-20.137	0.838	1.00	0.95	SEG1
ATOM	1617	CB	ALA	105	18.082	-21.692	0.012	1.00	0.98	SEG1
ATOM	1618	HB1	ALA	105	17.663	-21.821	0.999	1.00	1.41	SEG1
ATOM	1619	HB2	ALA	105	18.797	-22.477	-0.181	1.00	1.44	SEG1
ATOM	1620	HB3	ALA	105	17.291	-21.738	-0.723	1.00	1.39	SEG1
ATOM	1621	C	ALA	105	19.719	-20.287	-1.280	1.00	1.03	SEG1
ATOM	1622	O	ALA	105	20.914	-20.482	-1.160	1.00	1.11	SEG1
ATOM	1623	N	ALA	106	19.177	-20.032	-2.443	1.00	1.04	SEG1
ATOM	1624	HN	ALA	106	18.210	-19.882	-2.503	1.00	0.98	SEG1
ATOM	1625	CA	ALA	106	20.019	-19.972	-3.678	1.00	1.14	SEG1
ATOM	1626	HA	ALA	106	20.573	-20.889	-3.798	1.00	1.22	SEG1
ATOM	1627	CB	ALA	106	19.027	-19.812	-4.833	1.00	1.13	SEG1
ATOM	1628	HB1	ALA	106	19.566	-19.771	-5.767	1.00	1.31	SEG1
ATOM	1629	HB2	ALA	106	18.466	-18.898	-4.701	1.00	1.58	SEG1
ATOM	1630	HB3	ALA	106	18.349	-20.652	-4.844	1.00	1.56	SEG1
ATOM	1631	C	ALA	106	20.974	-18.775	-3.628	1.00	1.18	SEG1
ATOM	1632	O	ALA	106	22.086	-18.844	-4.117	1.00	1.29	SEG1
ATOM	1633	N	LEU	107	20.546	-17.677	-3.054	1.00	1.11	SEG1
ATOM	1634	HN	LEU	107	19.644	-17.642	-2.674	1.00	1.02	SEG1
ATOM	1635	CA	LEU	107	21.426	-16.473	-2.990	1.00	1.18	SEG1
ATOM	1636	HA	LEU	107	22.066	-16.433	-3.857	1.00	1.27	SEG1
ATOM	1637	CB	LEU	107	20.463	-15.285	-3.003	1.00	1.12	SEG1
ATOM	1638	HB1	LEU	107	21.007	-14.376	-2.794	1.00	1.19	SEG1
ATOM	1639	HB2	LEU	107	19.701	-15.431	-2.251	1.00	1.05	SEG1
ATOM	1640	CG	LEU	107	19.808	-15.180	-4.382	1.00	1.12	SEG1
ATOM	1641	HG	LEU	107	19.415	-16.145	-4.666	1.00	1.14	SEG1
ATOM	1642	CD1	LEU	107	18.669	-14.161	-4.334	1.00	1.21	SEG1
ATOM	1643	HD11	LEU	107	18.997	-13.278	-3.806	1.00	1.78	SEG1
ATOM	1644	HD12	LEU	107	17.821	-14.592	-3.822	1.00	1.60	SEG1
ATOM	1645	HD13	LEU	107	18.383	-13.893	-5.341	1.00	1.41	SEG1
ATOM	1646	CD2	LEU	107	20.851	-14.730	-5.407	1.00	1.24	SEG1
ATOM	1647	HD21	LEU	107	21.517	-14.011	-4.953	1.00	1.62	SEG1
ATOM	1648	HD22	LEU	107	20.353	-14.277	-6.252	1.00	1.68	SEG1
ATOM	1649	HD23	LEU	107	21.419	-15.585	-5.742	1.00	1.59	SEG1
ATOM	1650	C	LEU	107	22.270	-16.490	-1.708	1.00	1.21	SEG1
ATOM	1651	O	LEU	107	23.433	-16.849	-1.734	1.00	1.31	SEG1
ATOM	1652	N	ALA	108	21.701	-16.107	-0.589	1.00	1.13	SEG1
ATOM	1653	HN	ALA	108	20.762	-15.822	-0.590	1.00	1.05	SEG1
ATOM	1654	CA	ALA	108	22.483	-16.105	0.688	1.00	1.17	SEG1
ATOM	1655	HA	ALA	108	23.036	-17.024	0.790	1.00	1.21	SEG1
ATOM	1656	CB	ALA	108	23.449	-14.927	0.561	1.00	1.29	SEG1
ATOM	1657	HB1	ALA	108	24.163	-14.958	1.371	1.00	1.64	SEG1
ATOM	1658	HB2	ALA	108	22.896	-14.001	0.603	1.00	1.68	SEG1
ATOM	1659	HB3	ALA	108	23.972	-14.991	-0.382	1.00	1.65	SEG1
ATOM	1660	C	ALA	108	21.556	-15.907	1.891	1.00	1.08	SEG1
ATOM	1661	O	ALA	108	21.911	-15.258	2.857	1.00	1.13	SEG1
ATOM	1662	N	GLN	109	20.372	-16.458	1.836	1.00	0.97	SEG1
ATOM	1663	HN	GLN	109	20.112	-16.973	1.044	1.00	0.95	SEG1
ATOM	1664	CA	GLN	109	19.413	-16.303	2.971	1.00	0.90	SEG1
ATOM	1665	HA	GLN	109	19.905	-15.884	3.833	1.00	0.96	SEG1
ATOM	1666	CB	GLN	109	18.352	-15.334	2.447	1.00	0.85	SEG1
ATOM	1667	HB1	GLN	109	17.531	-15.286	3.146	1.00	0.80	SEG1
ATOM	1668	HB2	GLN	109	17.990	-15.680	1.489	1.00	0.82	SEG1
ATOM	1669	CG	GLN	109	18.965	-13.942	2.289	1.00	0.98	SEG1
ATOM	1670	HG1	GLN	109	19.887	-14.017	1.734	1.00	1.16	SEG1
ATOM	1671	HG2	GLN	109	19.165	-13.527	3.265	1.00	1.25	SEG1
ATOM	1672	CD	GLN	109	17.994	-13.028	1.537	1.00	1.42	SEG1
ATOM	1673	OE1	GLN	109	17.061	-13.490	0.910	1.00	1.91	SEG1
ATOM	1674	NE2	GLN	109	18.183	-11.738	1.565	1.00	1.74	SEG1
ATOM	1675	HE21	GLN	109	18.940	-11.366	2.064	1.00	1.72	SEG1
ATOM	1676	HE22	GLN	109	17.569	-11.142	1.087	1.00	2.24	SEG1
ATOM	1677	C	GLN	109	18.775	-17.652	3.311	1.00	0.85	SEG1
ATOM	1678	O	GLN	109	18.443	-18.423	2.432	1.00	0.83	SEG1
ATOM	1679	N	HIS	110	18.587	-17.941	4.576	1.00	0.87	SEG1
ATOM	1680	HN	HIS	110	18.852	-17.302	5.270	1.00	0.92	SEG1
ATOM	1681	CA	HIS	110	17.951	-19.240	4.950	1.00	0.89	SEG1

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ATOM	1682	HA	HIS	110	18.501	-20.067	4.529	1.00	0.94	SEG1
ATOM	1683	CB	HIS	110	17.997	-19.298	6.479	1.00	0.95	SEG1
ATOM	1684	HB1	HIS	110	17.393	-20.124	6.825	1.00	1.35	SEG1
ATOM	1685	HB2	HIS	110	17.611	-18.375	6.885	1.00	1.16	SEG1
ATOM	1686	CG	HIS	110	19.416	-19.492	5.934	1.00	1.24	SEG1
ATOM	1687	ND1	HIS	110	20.255	-18.426	7.216	1.00	2.20	SEG1
ATOM	1688	HD1	HIS	110	20.025	-17.476	7.152	1.00	2.77	SEG1
ATOM	1689	CD2	HIS	110	20.157	-20.624	7.162	1.00	1.69	SEG1
ATOM	1690	HD2	HIS	110	19.802	-21.636	7.034	1.00	2.16	SEG1
ATOM	1691	CE1	HIS	110	21.442	-18.934	7.595	1.00	2.66	SEG1
ATOM	1692	HE1	HIS	110	22.295	-18.335	7.877	1.00	3.53	SEG1
ATOM	1693	NE2	HIS	110	21.436	-20.270	7.580	1.00	2.33	SEG1
ATOM	1694	C	HIS	110	16.503	-19.253	4.460	1.00	0.83	SEG1
ATOM	1695	O	HIS	110	16.013	-20.249	3.962	1.00	1.08	SEG1
ATOM	1696	N	SEP	111	15.820	-18.141	4.587	1.00	0.68	SEG1
ATOM	1697	HN	SEP	111	16.249	-17.353	4.984	1.00	0.77	SEG1
ATOM	1698	CA	SER	111	14.400	-18.053	4.121	1.00	0.61	SEG1
ATOM	1699	HA	SER	111	14.329	-18.326	3.080	1.00	0.63	SEG1
ATOM	1700	CB	SER	111	13.618	-19.051	4.982	1.00	0.66	SEG1
ATOM	1701	HB1	SER	111	13.856	-18.888	6.025	1.00	1.14	SEG1
ATOM	1702	HB2	SER	111	13.889	-20.056	4.708	1.00	1.08	SEG1
ATOM	1703	CG	SER	111	12.225	-18.866	4.767	1.00	1.38	SEG1
ATOM	1704	HG	SER	111	11.755	-19.295	5.485	1.00	1.53	SEG1
ATOM	1705	C	SER	111	13.862	-16.637	4.341	1.00	0.51	SEG1
ATOM	1706	O	SER	111	13.860	-16.132	5.449	1.00	0.54	SEG1
ATOM	1707	N	VAL	112	13.392	-16.000	3.299	1.00	0.47	SEG1
ATOM	1708	HN	VAL	112	13.395	-16.433	2.420	1.00	0.51	SEG1
ATOM	1709	CA	VAL	112	12.834	-14.620	3.448	1.00	0.42	SEG1
ATOM	1710	HA	VAL	112	12.698	-14.386	4.492	1.00	0.41	SEG1
ATOM	1711	CB	VAL	112	13.882	-13.674	2.833	1.00	0.54	SEG1
ATOM	1712	HB	VAL	112	14.788	-13.721	3.420	1.00	0.61	SEG1
ATOM	1713	CG1	VAL	112	14.195	-14.094	1.390	1.00	0.61	SEG1
ATOM	1714	HG11	VAL	112	13.434	-14.775	1.038	1.00	1.32	SEG1
ATOM	1715	HG12	VAL	112	15.158	-14.584	1.361	1.00	1.02	SEG1
ATOM	1716	HG13	VAL	112	14.219	-13.221	0.754	1.00	1.21	SEG1
ATOM	1717	CG2	VAL	112	13.344	-12.234	2.844	1.00	0.58	SEG1
ATOM	1718	HG21	VAL	112	12.561	-12.147	3.587	1.00	1.18	SEG1
ATOM	1719	HG22	VAL	112	12.943	-11.993	1.870	1.00	1.21	SEG1
ATOM	1720	HG23	VAL	112	14.145	-11.552	3.082	1.00	1.13	SEG1
ATOM	1721	C	VAL	112	11.491	-14.519	2.704	1.00	0.40	SEG1
ATOM	1722	O	VAL	112	11.400	-13.850	1.693	1.00	0.46	SEG1
ATOM	1723	N	PRO	113	10.483	-15.181	3.228	1.00	0.41	SEG1
ATOM	1724	CA	PRO	113	9.151	-15.136	2.587	1.00	0.50	SEG1
ATOM	1725	HA	PRO	113	9.232	-15.296	1.525	1.00	0.57	SEG1
ATOM	1726	CB	PRO	113	8.398	-16.294	3.234	1.00	0.64	SEG1
ATOM	1727	HB1	PRO	113	8.486	-17.184	2.631	1.00	0.72	SEG1
ATOM	1728	HB2	PRO	113	7.357	-16.032	3.370	1.00	0.71	SEG1
ATOM	1729	CG	PRO	113	9.061	-16.507	4.558	1.00	0.62	SEG1
ATOM	1730	HG1	PRO	113	9.056	-17.557	4.807	1.00	0.75	SEG1
ATOM	1731	HG2	PRO	113	8.542	-15.944	5.321	1.00	0.66	SEG1
ATOM	1732	CD	PRO	113	10.483	-16.022	4.438	1.00	0.47	SEG1
ATOM	1733	HD2	PRO	113	10.757	-15.440	5.308	1.00	0.47	SEG1
ATOM	1734	HD1	PRO	113	11.158	-16.854	4.311	1.00	0.50	SEG1
ATOM	1735	C	PRO	113	8.458	-13.803	2.886	1.00	0.47	SEG1
ATOM	1736	O	PRO	113	7.929	-13.597	3.962	1.00	0.53	SEG1
ATOM	1737	N	LEU	114	8.447	-12.905	1.934	1.00	0.51	SEG1
ATOM	1738	HN	LEU	114	8.872	-13.102	1.073	1.00	0.59	SEG1
ATOM	1739	CA	LEU	114	7.778	-11.586	2.147	1.00	0.54	SEG1
ATOM	1740	HA	LEU	114	7.662	-11.378	3.199	1.00	0.50	SEG1
ATOM	1741	CB	LEU	114	8.709	-10.559	1.501	1.00	0.62	SEG1
ATOM	1742	HB1	LEU	114	8.194	-9.615	1.409	1.00	1.24	SEG1
ATOM	1743	HB2	LEU	114	9.001	-10.907	0.521	1.00	1.15	SEG1
ATOM	1744	CG	LEU	114	9.953	-10.375	2.368	1.00	1.00	SEG1
ATOM	1745	HC	LEU	114	10.362	-11.341	2.624	1.00	1.82	SEG1
ATOM	1746	CD1	LEU	114	10.995	-9.566	1.594	1.00	1.41	SEG1
ATOM	1747	HD11	LEU	114	10.713	-8.523	1.593	1.00	1.88	SEG1
ATOM	1748	HD12	LEU	114	11.047	-9.926	0.578	1.00	1.82	SEG1
ATOM	1749	HD13	LEU	114	11.960	-9.677	2.066	1.00	2.03	SEG1
ATOM	1750	CD2	LEU	114	9.572	-9.617	3.643	1.00	1.66	SEG1
ATOM	1751	HD21	LEU	114	8.980	-10.256	4.279	1.00	2.15	SEG1
ATOM	1752	HD22	LEU	114	8.998	-8.740	3.382	1.00	2.23	SEG1
ATOM	1753	HD23	LEU	114	10.469	-9.319	4.166	1.00	2.01	SEG1
ATOM	1754	C	LEU	114	6.427	-11.590	1.442	1.00	0.65	SEG1
ATOM	1755	O	LEU	114	6.353	-11.590	0.228	1.00	0.86	SEG1
ATOM	1756	N	GLN	115	5.359	-11.604	2.194	1.00	0.58	SEG1
ATOM	1757	HN	GLN	115	5.447	-11.613	3.170	1.00	0.51	SEG1
ATOM	1758	CA	GLN	115	4.008	-11.628	1.570	1.00	0.71	SEG1

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ATOM	1759	HA	GLN	115	4.095	-11.648	0.495	1.00	0.84	SEG1
ATOM	1760	CB	GLN	115	3.368	-12.929	2.062	1.00	0.86	SEG1
ATOM	1761	HB1	GLN	115	3.309	-12.918	3.139	1.00	0.94	SEG1
ATOM	1761	HB2	GLN	115	3.972	-13.768	1.742	1.00	1.29	SEG1
ATOM	1763	CG	GLN	115	1.959	-13.065	1.477	1.00	1.26	SEG1
ATOM	1764	HG1	GLN	115	2.018	-13.077	0.398	1.00	1.92	SEG1
ATOM	1765	HG2	GLN	115	1.358	-12.225	1.793	1.00	1.92	SEG1
ATOM	1766	CD	GLN	115	1.312	-14.366	1.963	1.00	0.81	SEG1
ATOM	1767	OE1	GLN	115	1.964	-15.205	2.553	1.00	1.12	SEG1
ATOM	1768	NE2	GLN	115	0.045	-14.571	1.730	1.00	0.81	SEG1
ATOM	1769	HE21	GLN	115	-0.481	-13.897	1.248	1.00	0.83	SEG1
ATOM	1770	HE22	GLN	115	-0.382	-15.399	2.035	1.00	1.18	SEG1
ATOM	1771	C	GLN	115	3.189	-10.414	2.011	1.00	0.56	SEG1
ATOM	1772	O	GLN	115	3.430	-9.822	3.053	1.00	0.46	SEG1
ATOM	1773	N	LEU	116	2.224	-10.047	1.210	1.00	0.59	SEG1
ATOM	1774	HN	LEU	116	2.068	-10.548	0.382	1.00	0.70	SEG1
ATOM	1775	CA	LEU	116	1.362	-8.878	1.533	1.00	0.48	SEG1
ATOM	1776	HA	LEU	116	1.742	-8.340	2.382	1.00	0.42	SEG1
ATOM	1777	CB	LEU	116	1.414	-7.993	0.284	1.00	0.53	SEG1
ATOM	1778	HB1	LEU	116	0.809	-7.113	0.443	1.00	0.52	SEG1
ATOM	1779	HB2	LEU	116	1.027	-8.545	-0.560	1.00	0.62	SEG1
ATOM	1780	CG	LEU	116	2.859	-7.567	-0.008	1.00	0.55	SEG1
ATOM	1781	HG	LEU	116	3.464	-8.446	-0.174	1.00	0.59	SEG1
ATOM	1782	CD1	LEU	116	2.880	-6.691	-1.261	1.00	0.69	SEG1
ATOM	1783	HD11	LEU	116	2.685	-5.665	-0.986	1.00	1.24	SEG1
ATOM	1784	HD12	LEU	116	2.119	-7.030	-1.950	1.00	1.29	SEG1
ATOM	1785	HD13	LEU	116	3.849	-6.760	-1.732	1.00	1.07	SEG1
ATOM	1786	CD2	LEU	116	3.422	-6.771	1.175	1.00	0.48	SEG1
ATOM	1787	HD21	LEU	116	2.704	-6.026	1.484	1.00	1.17	SEG1
ATOM	1788	HD22	LEU	116	4.339	-6.284	0.876	1.00	1.11	SEG1
ATOM	1789	HD23	LEU	116	3.623	-7.442	1.998	1.00	1.09	SEG1
ATOM	1790	C	LEU	116	-0.071	-9.349	1.783	1.00	0.48	SEG1
ATOM	1791	O	LEU	116	-0.511	-10.339	1.229	1.00	0.53	SEG1
ATOM	1792	N	GLU	117	-0.801	-8.643	2.604	1.00	0.43	SEG1
ATOM	1793	HN	GLU	117	-0.421	-7.846	3.030	1.00	0.40	SEG1
ATOM	1794	CA	GLU	117	-2.217	-9.030	2.884	1.00	0.44	SEG1
ATOM	1795	HA	GLU	117	-2.535	-9.813	2.214	1.00	0.49	SEG1
ATOM	1796	CB	GLU	117	-2.220	-9.533	4.327	1.00	0.47	SEG1
ATOM	1797	HB1	GLU	117	-3.228	-9.790	4.617	1.00	0.74	SEG1
ATOM	1798	HB2	GLU	117	-1.844	-8.760	4.979	1.00	0.64	SEG1
ATOM	1799	CG	GLU	117	-1.329	-10.776	4.433	1.00	0.95	SEG1
ATOM	1800	HG1	GLU	117	-0.322	-10.520	4.137	1.00	1.24	SEG1
ATOM	1801	HG2	GLU	117	-1.709	-11.544	3.775	1.00	1.29	SEG1
ATOM	1802	CD	GLU	117	-1.314	-11.304	5.875	1.00	0.99	SEG1
ATOM	1803	OE1	GLU	117	-1.813	-10.617	6.754	1.00	1.61	SEG1
ATOM	1804	OE2	GLU	117	-0.799	-12.392	6.075	1.00	1.23	SEG1
ATOM	1805	C	GLU	117	-3.104	-7.796	2.731	1.00	0.37	SEG1
ATOM	1806	O	GLU	117	-2.623	-6.682	2.781	1.00	0.39	SEG1
ATOM	1807	N	LEU	118	-4.386	-7.975	2.527	1.00	0.34	SEG1
ATOM	1808	HN	LEU	118	-4.755	-8.881	2.477	1.00	0.37	SEG1
ATOM	1809	CA	LEU	118	-5.277	-6.788	2.355	1.00	0.29	SEG1
ATOM	1810	HA	LEU	118	-4.699	-5.878	2.367	1.00	0.27	SEG1
ATOM	1811	CB	LEU	118	-5.931	-6.962	0.987	1.00	0.30	SEG1
ATOM	1812	HB1	LEU	118	-6.637	-6.161	0.826	1.00	0.28	SEG1
ATOM	1813	HB2	LEU	118	-6.451	-7.907	0.956	1.00	0.32	SEG1
ATOM	1814	CG	LEU	118	-4.865	-6.918	-0.104	1.00	0.31	SEG1
ATOM	1815	HG	LEU	118	-4.058	-7.590	0.150	1.00	0.34	SEG1
ATOM	1816	CD1	LEU	118	-5.483	-7.338	-1.438	1.00	0.36	SEG1
ATOM	1817	HD11	LEU	118	-4.712	-7.729	-2.085	1.00	1.00	SEG1
ATOM	1818	HD12	LEU	118	-5.946	-6.481	-1.906	1.00	1.02	SEG1
ATOM	1819	HD13	LEU	118	-6.229	-8.099	-1.265	1.00	1.09	SEG1
ATOM	1820	CD2	LEU	118	-4.329	-5.491	-0.218	1.00	0.29	SEG1
ATOM	1821	HD21	LEU	118	-3.997	-5.309	-1.229	1.00	1.02	SEG1
ATOM	1822	HD22	LEU	118	-3.502	-5.366	0.461	1.00	1.06	SEG1
ATOM	1823	HD23	LEU	118	-5.113	-4.790	0.036	1.00	1.07	SEG1
ATOM	1824	C	LEU	118	-6.364	-6.741	3.418	1.00	0.32	SEG1
ATOM	1825	O	LEU	118	-7.051	-7.715	3.665	1.00	0.38	SEG1
ATOM	1826	N	ARG	119	-6.556	-5.595	4.009	1.00	0.29	SEG1
ATOM	1827	HN	ARG	119	-6.006	-4.816	3.761	1.00	0.28	SEG1
ATOM	1828	CA	ARG	119	-7.638	-5.447	5.019	1.00	0.34	SEG1
ATOM	1829	HA	ARG	119	-8.268	-6.324	5.027	1.00	0.38	SEG1
ATOM	1830	CB	ARG	119	-6.939	-5.284	6.366	1.00	0.38	SEG1
ATOM	1831	HB1	ARG	119	-6.351	-4.379	6.361	1.00	0.61	SEG1
ATOM	1832	HB2	ARG	119	-6.294	-6.134	6.543	1.00	0.77	SEG1
ATOM	1833	CG	ARG	119	-7.993	-5.200	7.473	1.00	0.70	SEG1
ATOM	1834	HG1	ARG	119	-8.644	-6.058	7.417	1.00	1.23	SEG1
ATOM	1835	HG2	ARG	119	-8.574	-4.297	7.348	1.00	1.36	SEG1

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ATOM	1836	CD	ARG	119	-7.300	-5.177	8.835	1.00	0.84	SEG1
ATOM	1837	HD1	ARG	119	-6.660	-4.314	8.916	1.00	1.40	SEG1
ATOM	1838	HD2	ARG	119	-6.730	-6.086	8.975	1.00	1.09	SEG1
ATOM	1839	NE	ARG	119	-8.402	-5.094	9.846	1.00	1.43	SEG1
ATOM	1840	HE	ARG	119	-9.323	-4.949	9.545	1.00	1.78	SEG1
ATOM	1841	CZ	ARG	119	-8.156	-5.211	11.137	1.00	2.13	SEG1
ATOM	1842	NH1	ARG	119	-9.146	-5.133	11.984	1.00	2.72	SEG1
ATOM	1843	HH11	ARG	119	-10.078	-4.987	11.652	1.00	2.79	SEG1
ATOM	1844	HH12	ARG	119	-8.973	-5.220	12.964	1.00	3.33	SEG1
ATOM	1845	NH2	ARG	119	-6.938	-5.401	11.591	1.00	2.69	SEG1
ATOM	1846	HH21	ARG	119	-6.165	-5.461	10.961	1.00	2.58	SEG1
ATOM	1847	HH22	ARG	119	-6.783	-5.485	12.576	1.00	3.46	SEG1
ATOM	1848	C	ARG	119	-8.453	-4.207	4.667	1.00	0.31	SEG1
ATOM	1849	O	ARG	119	-8.041	-3.090	4.921	1.00	0.30	SEG1
ATOM	1850	N	ALA	120	-9.593	-4.399	4.059	1.00	0.33	SEG1
ATOM	1851	HN	ALA	120	-9.885	-5.314	3.846	1.00	0.34	SEG1
ATOM	1852	CA	ALA	120	-10.432	-3.231	3.658	1.00	0.33	SEG1
ATOM	1853	HA	ALA	120	-9.842	-2.327	3.679	1.00	0.33	SEG1
ATOM	1854	CB	ALA	120	-10.859	-3.532	2.221	1.00	0.35	SEG1
ATOM	1855	HB1	ALA	120	-11.394	-4.469	2.194	1.00	1.06	SEG1
ATOM	1856	HB2	ALA	120	-9.984	-3.598	1.592	1.00	1.10	SEG1
ATOM	1857	HB3	ALA	120	-11.501	-2.740	1.863	1.00	1.05	SEG1
ATOM	1858	C	ALA	120	-11.665	-3.096	4.562	1.00	0.38	SEG1
ATOM	1859	O	ALA	120	-12.261	-2.038	4.649	1.00	0.42	SEG1
ATOM	1860	N	GLY	121	-12.054	-4.154	5.230	1.00	0.40	SEG1
ATOM	1861	HN	GLY	121	-11.562	-4.996	5.142	1.00	0.40	SEG1
ATOM	1862	CA	GLY	121	-13.250	-4.083	6.122	1.00	0.46	SEG1
ATOM	1863	HA1	GLY	121	-14.119	-3.816	5.540	1.00	0.49	SEG1
ATOM	1864	HA2	GLY	121	-13.085	-3.338	6.887	1.00	0.48	SEG1
ATOM	1865	C	GLY	121	-13.481	-5.446	6.776	1.00	0.49	SEG1
ATOM	1866	O	GLY	121	-12.594	-6.003	7.396	1.00	0.50	SEG1
ATOM	1867	N	ALA	122	-14.666	-5.986	6.640	1.00	0.53	SEG1
ATOM	1868	HN	ALA	122	-15.361	-5.515	6.134	1.00	0.54	SEG1
ATOM	1869	CA	ALA	122	-14.962	-7.317	7.252	1.00	0.58	SEG1
ATOM	1870	HA	ALA	122	-14.535	-7.376	8.241	1.00	0.60	SEG1
ATOM	1871	CB	ALA	122	-16.487	-7.382	7.341	1.00	0.64	SEG1
ATOM	1872	HB1	ALA	122	-16.917	-7.030	6.416	1.00	1.23	SEG1
ATOM	1873	HB2	ALA	122	-16.826	-6.759	8.156	1.00	1.17	SEG1
ATOM	1874	HB3	ALA	122	-16.794	-8.402	7.516	1.00	1.23	SEG1
ATOM	1875	C	ALA	122	-14.425	-8.458	6.372	1.00	0.56	SEG1
ATOM	1876	O	ALA	122	-14.455	-9.610	6.763	1.00	0.60	SEG1
ATOM	1877	N	GLU	123	-13.932	-8.151	5.193	1.00	0.52	SEG1
ATOM	1878	HN	GLU	123	-13.912	-7.220	4.894	1.00	0.51	SEG1
ATOM	1879	CA	GLU	123	-13.392	-9.220	4.301	1.00	0.51	SEG1
ATOM	1880	HA	GLU	123	-13.761	-10.187	4.606	1.00	0.56	SEG1
ATOM	1881	CB	GLU	123	-13.919	-8.875	2.908	1.00	0.52	SEG1
ATOM	1882	HB1	GLU	123	-13.430	-9.494	2.173	1.00	0.60	SEG1
ATOM	1883	HB2	GLU	123	-13.716	-7.835	2.696	1.00	0.62	SEG1
ATOM	1884	CG	GLU	123	-15.432	-9.122	2.860	1.00	0.81	SEG1
ATOM	1885	HG1	GLU	123	-15.919	-8.503	3.598	1.00	1.14	SEG1
ATOM	1886	HG2	GLU	123	-15.630	-10.161	3.078	1.00	1.08	SEG1
ATOM	1887	CD	GLU	123	-15.985	-8.781	1.468	1.00	1.04	SEG1
ATOM	1888	OE1	GLU	123	-15.256	-8.207	0.673	1.00	1.61	SEG1
ATOM	1889	OE2	GLU	123	-17.134	-9.109	1.219	1.00	1.62	SEG1
ATOM	1890	C	GLU	123	-11.861	-9.201	4.324	1.00	0.46	SEG1
ATOM	1891	O	GLU	123	-11.242	-8.169	4.146	1.00	0.45	SEG1
ATOM	1892	N	ARG	124	-11.251	-10.338	4.545	1.00	0.49	SEG1
ATOM	1893	HN	ARG	124	-11.777	-11.152	4.688	1.00	0.54	SEG1
ATOM	1894	CA	ARG	124	-9.759	-10.401	4.587	1.00	0.49	SEG1
ATOM	1895	HA	ARG	124	-9.364	-9.483	4.993	1.00	0.48	SEG1
ATOM	1896	CB	ARG	124	-9.435	-11.562	5.529	1.00	0.56	SEG1
ATOM	1897	HB1	ARG	124	-8.385	-11.798	5.467	1.00	0.58	SEG1
ATOM	1898	HB2	ARG	124	-10.018	-12.427	5.248	1.00	0.57	SEG1
ATOM	1899	CG	ARG	124	-9.781	-11.156	6.954	1.00	0.61	SEG1
ATOM	1900	HG1	ARG	124	-10.832	-10.919	7.028	1.00	0.61	SEG1
ATOM	1901	HG2	ARG	124	-9.199	-10.288	7.240	1.00	0.60	SEG1
ATOM	1902	CD	ARG	124	-9.461	-12.312	7.918	1.00	0.70	SEG1
ATOM	1903	HD1	ARG	124	-8.407	-12.546	7.886	1.00	1.17	SEG1
ATOM	1904	HD2	ARG	124	-10.048	-13.182	7.661	1.00	1.17	SEG1
ATOM	1905	NE	ARG	124	-9.832	-11.814	9.281	1.00	1.59	SEG1
ATOM	1906	HE	ARG	124	-10.282	-10.947	9.372	1.00	2.17	SEG1
ATOM	1907	CZ	ARG	124	-9.567	-12.517	10.365	1.00	2.20	SEG1
ATOM	1908	NH1	ARG	124	-9.908	-12.042	11.532	1.00	3.24	SEG1
ATOM	1909	HH11	ARG	124	-10.364	-11.155	11.596	1.00	3.68	SEG1
ATOM	1910	HH12	ARG	124	-9.713	-12.565	12.361	1.00	3.76	SEG1
ATOM	1911	NH2	ARG	124	-8.967	-13.683	10.300	1.00	2.24	SEG1
ATOM	1912	HH21	ARG	124	-8.696	-14.065	9.418	1.00	2.00	SEG1

FIG. 2 (25 of 35)

ATC4	1913	HR23	ARG	124	-8.780	-14.194	11.139	1.00	2.90	SEG1
ATC4	1914	C	ARG	124	-9.185	-10.648	3.183	1.00	0.46	SEG1
ATC4	1915	O	ARG	124	-9.872	-10.511	2.193	1.00	0.44	SEG1
ATC4	1916	N	LEU	125	-7.923	-10.993	3.139	1.00	0.49	SEG1
ATC4	1917	HN	LEU	125	-7.393	-11.079	4.928	1.00	0.53	SEG1
ATC4	1918	CA	LEU	125	-7.283	-11.230	1.777	1.00	0.49	SEG1
ATC4	1919	HA	LEU	125	-7.297	-10.325	1.191	1.00	0.50	SEG1
ATC4	1920	CB	LEU	125	-5.935	-11.627	2.096	1.00	0.58	SEG1
ATC4	1921	HB1	LEU	125	-5.834	-12.518	2.794	1.00	0.60	SEG1
ATC4	1922	HB2	LEU	125	-5.353	-10.823	2.635	1.00	0.62	SEG1
ATC4	1923	CG	LEU	125	-5.069	-11.899	0.798	1.00	0.62	SEG1
ATC4	1924	HG	LEU	125	-5.615	-12.621	0.205	1.00	0.79	SEG1
ATC4	1925	CD1	LEU	125	-4.915	-10.600	-0.003	1.00	1.11	SEG1
ATC4	1926	HD11	LEU	125	-3.896	-10.254	0.051	1.00	1.61	SEG1
ATC4	1927	HD12	LEU	125	-5.573	-9.844	0.397	1.00	1.75	SEG1
ATC4	1928	HD13	LEU	125	-5.167	-10.784	-1.037	1.00	1.55	SEG1
ATC4	1929	CD2	LEU	125	-3.686	-12.463	1.119	1.00	1.28	SEG1
ATC4	1930	HD21	LEU	125	-3.790	-13.260	1.959	1.00	1.80	SEG1
ATC4	1931	HD22	LEU	125	-3.071	-11.679	1.556	1.00	1.84	SEG1
ATC4	1932	HD23	LEU	125	-3.222	-12.847	0.242	1.00	1.73	SEG1
ATC4	1933	C	LEU	125	-7.294	-12.362	1.008	1.00	0.46	SEG1
ATC4	1934	O	LEU	125	-8.299	-12.236	-0.144	1.00	0.45	SEG1
ATC4	1935	N	ASP	126	-8.261	-13.458	1.688	1.00	0.49	SEG1
ATC4	1936	HN	ASP	126	-8.007	-13.537	2.632	1.00	0.53	SEG1
ATC4	1937	CA	ASP	126	-8.956	-14.590	1.001	1.00	0.50	SEG1
ATC4	1938	HA	ASP	126	-8.381	-14.913	0.145	1.00	0.49	SEG1
ATC4	1939	CB	ASP	126	-9.025	-15.723	2.038	1.00	0.57	SEG1
ATC4	1940	HB1	ASP	126	-8.025	-15.964	2.369	1.00	0.58	SEG1
ATC4	1941	HB2	ASP	126	-9.469	-16.596	1.583	1.00	0.59	SEG1
ATC4	1942	CG	ASP	126	-9.868	-15.299	3.251	1.00	0.62	SEG1
ATC4	1943	OD1	ASP	126	-10.047	-14.108	3.446	1.00	1.28	SEG1
ATC4	1944	OD2	ASP	126	-10.322	-16.179	3.964	1.00	1.21	SEG1
ATC4	1945	C	ASP	126	-10.360	-14.158	0.559	1.00	0.51	SEG1
ATC4	1946	O	ASP	126	-10.948	-14.746	-0.329	1.00	0.53	SEG1
ATC4	1947	N	ALA	127	-10.899	-13.135	1.177	1.00	0.52	SEG1
ATC4	1948	HN	ALA	127	-10.406	-12.680	1.890	1.00	0.52	SEG1
ATC4	1949	CA	ALA	127	-12.263	-12.658	0.803	1.00	0.57	SEG1
ATC4	1950	HA	ALA	127	-12.905	-13.496	0.584	1.00	0.60	SEG1
ATC4	1951	CB	ALA	127	-12.773	-11.927	2.046	1.00	0.64	SEG1
ATC4	1952	HB1	ALA	127	-12.036	-11.203	2.368	1.00	1.19	SEG1
ATC4	1953	HB2	ALA	127	-12.944	-12.642	2.838	1.00	1.26	SEG1
ATC4	1954	HB3	ALA	127	-13.697	-11.422	1.811	1.00	1.17	SEG1
ATC4	1955	C	ALA	127	-12.209	-11.698	-0.398	1.00	0.55	SEG1
ATC4	1956	O	ALA	127	-13.236	-11.313	-0.925	1.00	0.62	SEG1
ATC4	1957	N	LEU	128	-11.031	-11.301	-0.832	1.00	0.50	SEG1
ATC4	1958	HN	LEU	128	-10.212	-11.614	-0.396	1.00	0.47	SEG1
ATC4	1959	CA	LEU	128	-10.943	-10.360	-1.994	1.00	0.52	SEG1
ATC4	1960	HA	LEU	128	-11.895	-9.885	-2.163	1.00	0.57	SEG1
ATC4	1961	CB	LEU	128	-9.903	-9.313	-1.585	1.00	0.53	SEG1
ATC4	1962	HB1	LEU	128	-9.780	-8.598	-2.385	1.00	0.87	SEG1
ATC4	1963	HB2	LEU	128	-8.958	-9.801	-1.393	1.00	0.70	SEG1
ATC4	1964	CG	LEU	128	-10.366	-8.583	-0.321	1.00	1.03	SEG1
ATC4	1965	HG	LEU	128	-10.624	-9.310	0.437	1.00	1.71	SEG1
ATC4	1966	CD1	LEU	128	-9.231	-7.694	0.196	1.00	1.29	SEG1
ATC4	1967	HD11	LEU	128	-8.668	-7.281	-0.641	1.00	1.79	SEG1
ATC4	1968	HD12	LEU	128	-8.562	-8.283	0.805	1.00	1.73	SEG1
ATC4	1969	HD13	LEU	128	-9.644	-6.891	0.789	1.00	1.77	SEG1
ATC4	1970	CD2	LEU	128	-11.594	-7.715	-0.640	1.00	1.58	SEG1
ATC4	1971	HD21	LEU	128	-11.309	-6.673	-0.659	1.00	2.06	SEG1
ATC4	1972	HD22	LEU	128	-12.346	-7.866	0.120	1.00	1.98	SEG1
ATC4	1973	HD23	LEU	128	-11.997	-7.994	-1.602	1.00	2.07	SEG1
ATC4	1974	C	LEU	128	-10.481	-11.091	-3.259	1.00	0.50	SEG1
ATC4	1975	O	LEU	128	-10.738	-10.647	-4.362	1.00	0.55	SEG1
ATC4	1976	N	LEU	129	-9.800	-12.201	-3.113	1.00	0.47	SEG1
ATC4	1977	HN	LEU	129	-9.601	-12.539	-2.214	1.00	0.46	SEG1
ATC4	1978	CA	LEU	129	-9.322	-12.948	-4.318	1.00	0.49	SEG1
ATC4	1979	HA	LEU	129	-8.785	-12.283	-4.976	1.00	0.53	SEG1
ATC4	1980	CB	LEU	129	-8.372	-14.026	-3.786	1.00	0.49	SEG1
ATC4	1981	HB1	LEU	129	-8.025	-14.635	-4.607	1.00	0.53	SEG1
ATC4	1982	HB2	LEU	129	-8.898	-14.647	-3.075	1.00	0.48	SEG1
ATC4	1983	CG	LEU	129	-7.167	-13.372	-3.095	1.00	0.50	SEG1
ATC4	1984	HG	LEU	129	-7.513	-12.769	-2.268	1.00	0.48	SEG1
ATC4	1985	CD1	LEU	129	-6.229	-14.460	-2.570	1.00	0.54	SEG1
ATC4	1986	HD11	LEU	129	-6.789	-15.368	-2.400	1.00	1.15	SEG1
ATC4	1987	HD12	LEU	129	-5.782	-14.134	-1.643	1.00	1.15	SEG1
ATC4	1988	HD13	LEU	129	-5.452	-14.648	-3.297	1.00	1.18	SEG1
ATC4	1989	CD2	LEU	129	-6.406	-12.490	-4.094	1.00	0.59	SEG1

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ATCM	1991	HD11	LEU	129	-6.990	-11.598	-4.296	1.00	1.13	SEG1
ATCM	1991	HD12	LEU	129	-6.253	-13.036	-5.013	1.00	1.11	SEG1
ATCM	1991	HD13	LEU	129	-5.450	-12.215	-3.675	1.00	1.28	SEG1
ATCM	1991	C	LEU	129	-10.501	-13.585	-5.061	1.00	0.54	SEG1
ATCM	1994	C	LEU	129	-10.542	-13.597	-5.278	1.00	0.60	SEG1
ATCM	1995	N	ALA	130	-11.454	-14.118	-4.340	1.00	0.56	SEG1
ATCM	1996	HN	ALA	130	-11.393	-14.098	-3.261	1.00	0.54	SEG1
ATCM	1997	CA	ALA	130	-12.631	-14.765	-5.003	1.00	0.64	SEG1
ATCM	1998	HA	ALA	130	-12.299	-15.512	-5.706	1.00	0.73	SEG1
ATCM	1999	CB	ALA	130	-13.418	-15.429	-3.871	1.00	0.72	SEG1
ATCM	2000	HB1	ALA	130	-14.057	-16.197	-4.281	1.00	1.23	SEG1
ATCM	2001	HB2	ALA	130	-14.022	-14.688	-3.369	1.00	1.22	SEG1
ATCM	2002	HB3	ALA	130	-12.730	-15.871	-3.166	1.00	1.30	SEG1
ATCM	2003	C	ALA	130	-13.497	-13.718	-5.711	1.00	0.59	SEG1
ATCM	2004	O	ALA	130	-13.759	-13.818	-6.895	1.00	0.64	SEG1
ATCM	2005	N	ASP	131	-13.948	-12.720	-4.992	1.00	0.55	SEG1
ATCM	2006	HN	ASP	131	-13.727	-12.667	-4.039	1.00	0.57	SEG1
ATCM	2007	CA	ASP	131	-14.807	-11.669	-5.616	1.00	0.53	SEG1
ATCM	2008	HA	ASP	131	-15.173	-12.001	-6.574	1.00	0.58	SEG1
ATCM	2009	CB	ASP	131	-15.975	-11.489	-4.646	1.00	0.61	SEG1
ATCM	2010	HB1	ASP	131	-16.552	-10.623	-4.934	1.00	0.74	SEG1
ATCM	2011	HB2	ASP	131	-15.592	-11.350	-3.645	1.00	0.91	SEG1
ATCM	2012	CG	ASP	131	-16.867	-12.730	-4.685	1.00	0.99	SEG1
ATCM	2013	OD1	ASP	131	-17.317	-13.145	-3.630	1.00	1.79	SEG1
ATCM	2014	OD2	ASP	131	-17.089	-13.242	-5.770	1.00	1.43	SEG1
ATCM	2015	C	ASP	131	-14.029	-10.359	-5.764	1.00	0.45	SEG1
ATCM	2016	O	ASP	131	-13.505	-9.828	-4.803	1.00	0.42	SEG1
ATCM	2017	N	GLU	132	-13.956	-9.838	-6.962	1.00	0.46	SEG1
ATCM	2018	HN	GLU	132	-14.390	-10.287	-7.717	1.00	0.51	SEG1
ATCM	2019	CA	GLU	132	-13.217	-8.561	-7.183	1.00	0.45	SEG1
ATCM	2020	HA	GLU	132	-12.367	-8.499	-6.522	1.00	0.44	SEG1
ATCM	2021	CB	GLU	132	-12.744	-8.621	-8.637	1.00	0.54	SEG1
ATCM	2022	HB1	GLU	132	-12.350	-7.659	-8.928	1.00	1.06	SEG1
ATCM	2023	HB2	GLU	132	-13.576	-8.878	-9.276	1.00	0.88	SEG1
ATCM	2024	CG	GLU	132	-11.645	-9.683	-8.771	1.00	1.23	SEG1
ATCM	2025	HG1	GLU	132	-12.041	-10.644	-8.476	1.00	1.67	SEG1
ATCM	2026	HG2	GLU	132	-10.818	-9.425	-8.126	1.00	1.77	SEG1
ATCM	2027	CD	GLU	132	-11.152	-9.764	-10.224	1.00	1.29	SEG1
ATCM	2028	OE1	GLU	132	-10.154	-10.430	-10.447	1.00	1.79	SEG1
ATCM	2029	OE2	GLU	132	-11.777	-9.167	-11.088	1.00	1.59	SEG1
ATCM	2030	C	GLU	132	-14.150	-7.366	-6.964	1.00	0.43	SEG1
ATCM	2031	O	GLU	132	-13.732	-6.321	-6.510	1.00	0.42	SEG1
ATCM	2032	N	GLU	133	-15.410	-7.511	-7.280	1.00	0.46	SEG1
ATCM	2033	HN	GLU	133	-15.727	-8.365	-7.645	1.00	0.49	SEG1
ATCM	2034	CA	GLU	133	-16.367	-6.377	-7.082	1.00	0.47	SEG1
ATCM	2035	HA	GLU	133	-16.061	-5.520	-7.660	1.00	0.50	SEG1
ATCM	2036	CB	GLU	133	-17.717	-6.894	-7.586	1.00	0.53	SEG1
ATCM	2037	HB1	GLU	133	-18.493	-6.195	-7.315	1.00	0.71	SEG1
ATCM	2038	HB2	GLU	133	-17.924	-7.855	-7.136	1.00	0.75	SEG1
ATCM	2039	CG	GLU	133	-17.676	-7.041	-9.111	1.00	0.96	SEG1
ATCM	2040	HG1	GLU	133	-16.898	-7.739	-9.381	1.00	1.39	SEG1
ATCM	2041	HG2	GLU	133	-17.465	-6.081	-9.558	1.00	1.25	SEG1
ATCM	2042	CD	GLU	133	-19.025	-7.560	-9.633	1.00	1.18	SEG1
ATCM	2043	OE1	GLU	133	-19.853	-7.956	-8.824	1.00	1.50	SEG1
ATCM	2044	OE2	GLU	133	-19.207	-7.554	-10.840	1.00	1.83	SEG1
ATCM	2045	C	GLU	133	-16.460	-6.017	-5.594	1.00	0.41	SEG1
ATCM	2046	O	GLU	133	-16.538	-4.858	-5.228	1.00	0.41	SEG1
ATCM	2047	N	ARG	134	-16.469	-7.008	-4.738	1.00	0.40	SEG1
ATCM	2048	HN	ARG	134	-16.419	-7.931	-5.065	1.00	0.42	SEG1
ATCM	2049	CA	ARG	134	-16.581	-6.744	-3.268	1.00	0.39	SEG1
ATCM	2050	HA	ARG	134	-17.502	-6.226	-3.058	1.00	0.42	SEG1
ATCM	2051	CB	ARG	134	-16.619	-8.118	-2.603	1.00	0.44	SEG1
ATCM	2052	HB1	ARG	134	-16.526	-8.002	-1.534	1.00	0.68	SEG1
ATCM	2053	HB2	ARG	134	-15.802	-8.720	-2.974	1.00	0.63	SEG1
ATCM	2054	CG	ARG	134	-17.947	-8.802	-2.925	1.00	0.71	SEG1
ATCM	2055	HG1	ARG	134	-18.030	-8.947	-3.991	1.00	1.28	SEG1
ATCM	2056	HG2	ARG	134	-18.764	-8.186	-2.578	1.00	1.12	SEG1
ATCM	2057	CD	ARG	134	-17.992	-10.158	-2.222	1.00	1.16	SEG1
ATCM	2058	HD1	ARG	134	-17.945	-10.023	-1.153	1.00	1.47	SEG1
ATCM	2059	HD2	ARG	134	-17.175	-10.779	-2.559	1.00	1.71	SEG1
ATCM	2060	NE	ARG	134	-19.306	-10.767	-2.603	1.00	1.51	SEG1
ATCM	2061	HE	ARG	134	-19.918	-10.269	-3.136	1.00	1.74	SEG1
ATCM	2062	CE	ARG	134	-19.655	-11.965	-2.177	1.00	2.12	SEG1
ATCM	2063	NH1	ARG	134	-20.802	-12.464	-2.549	1.00	2.52	SEG1
ATCM	2064	HH11	ARG	134	-21.405	-11.938	-3.149	1.00	2.52	SEG1
ATCM	2065	HH12	ARG	134	-21.078	-13.372	-2.234	1.00	3.06	SEG1
ATCM	2066	NH1	ARG	134	-18.878	-12.667	-1.383	1.00	2.76	SEG1

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ATOM	2067	RH21	ARG	134	-17.998	-12.302	-1.083	1.00	2.75	SEG1
ATOM	2068	RH22	ARG	134	-19.169	-13.573	-1.077	1.00	3.44	SEG1
ATOM	2069	C	ARG	134	-15.395	-5.931	-2.734	1.00	0.34	SEG1
ATOM	2070	O	ARG	134	-15.571	-5.067	-1.895	1.00	0.36	SEG1
ATOM	2071	N	CYS	135	-14.188	-6.202	-3.184	1.00	0.733	SEG1
ATOM	2072	HN	CYS	135	-14.056	-6.911	-3.848	1.00	0.34	SEG1
ATOM	2073	CA	CYS	135	-13.012	-5.435	-2.535	1.00	0.31	SEG1
ATOM	2074	HA	CYS	135	-12.925	-5.815	-1.534	1.00	0.32	SEG1
ATOM	2075	CB	CYS	135	-11.763	-6.017	-3.370	1.00	0.35	SEG1
ATOM	2076	HB1	CYS	135	-11.888	-7.086	-3.473	1.00	0.38	SEG1
ATOM	2077	HB2	CYS	135	-10.893	-5.826	-2.763	1.00	0.37	SEG1
ATOM	2078	SG	CYS	135	-11.491	-5.291	-5.018	1.00	0.41	SEG1
ATOM	2079	HG	CYS	135	-10.551	-5.117	-5.113	1.00	0.98	SEG1
ATOM	2080	C	CYS	135	-13.203	-3.927	-2.892	1.00	0.31	SEG1
ATOM	2081	G	CYS	135	-12.861	-3.112	-2.056	1.00	0.32	SEG1
ATOM	2082	N	LEU	136	-13.754	-3.560	-4.022	1.00	0.34	SEG1
ATOM	2083	HN	LEU	136	-14.026	-4.240	-4.675	1.00	0.36	SEG1
ATOM	2084	CA	LEU	136	-13.977	-2.111	-4.312	1.00	0.39	SEG1
ATOM	2085	HA	LEU	136	-13.043	-1.573	-4.293	1.00	0.41	SEG1
ATOM	2086	CB	LEU	136	-14.578	-2.074	-5.720	1.00	0.44	SEG1
ATOM	2087	HB1	LEU	136	-14.869	-1.063	-5.962	1.00	0.48	SEG1
ATOM	2088	HB2	LEU	136	-15.445	-2.718	-5.757	1.00	0.44	SEG1
ATOM	2089	CG	LEU	136	-13.538	-2.559	-6.730	1.00	0.46	SEG1
ATOM	2090	HG	LEU	136	-13.158	-3.522	-6.420	1.00	0.44	SEG1
ATOM	2091	CD1	LEU	136	-14.188	-2.690	-8.109	1.00	0.54	SEG1
ATOM	2092	HD11	LEU	136	-14.407	-1.707	-8.498	1.00	1.21	SEG1
ATOM	2093	HD12	LEU	136	-15.104	-3.256	-8.022	1.00	1.19	SEG1
ATOM	2094	HD13	LEU	136	-13.511	-3.201	-8.778	1.00	1.03	SEG1
ATOM	2095	CD2	LEU	136	-12.388	-1.553	-6.802	1.00	0.49	SEG1
ATOM	2096	HD21	LEU	136	-11.833	-1.575	-5.875	1.00	1.08	SEG1
ATOM	2097	HD22	LEU	136	-12.785	-0.562	-6.960	1.00	1.12	SEG1
ATOM	2098	HD23	LEU	136	-11.732	-1.813	-7.620	1.00	1.19	SEG1
ATOM	2099	C	LEU	136	-14.959	-1.519	-3.296	1.00	0.39	SEG1
ATOM	2100	O	LEU	136	-14.768	-0.426	-2.797	1.00	0.43	SEG1
ATOM	2101	N	SER	137	-16.009	-2.240	-2.989	1.00	0.38	SEG1
ATOM	2102	HN	SER	137	-16.134	-3.118	-3.410	1.00	0.37	SEG1
ATOM	2103	CA	SER	137	-17.017	-1.733	-2.003	1.00	0.43	SEG1
ATOM	2104	HA	SER	137	-17.464	-0.822	-2.363	1.00	0.48	SEG1
ATOM	2105	CB	SER	137	-18.078	-2.830	-1.911	1.00	0.48	SEG1
ATOM	2106	HB1	SER	137	-18.773	-2.591	-1.117	1.00	1.02	SEG1
ATOM	2107	HB2	SER	137	-17.605	-3.775	-1.658	1.00	1.21	SEG1
ATOM	2108	OG	SER	137	-18.768	-2.922	-3.150	1.00	1.35	SEG1
ATOM	2109	HG	SER	137	-18.239	-3.459	-3.745	1.00	1.57	SEG1
ATOM	2110	C	SER	137	-16.371	-1.502	-0.633	1.00	0.41	SEG1
ATOM	2111	O	SER	137	-16.674	-0.542	0.050	1.00	0.46	SEG1
ATOM	2112	N	CYS	138	-15.499	-2.386	-0.225	1.00	0.36	SEG1
ATOM	2113	HN	CYS	138	-15.285	-3.154	-0.795	1.00	0.35	SEG1
ATOM	2114	CA	CYS	138	-14.839	-2.239	1.111	1.00	0.37	SEG1
ATOM	2115	HA	CYS	138	-15.576	-2.070	1.877	1.00	0.42	SEG1
ATOM	2116	CB	CYS	138	-14.131	-3.571	1.358	1.00	0.39	SEG1
ATOM	2117	HB1	CYS	138	-13.360	-3.435	2.101	1.00	0.40	SEG1
ATOM	2118	HB2	CYS	138	-13.685	-3.917	0.436	1.00	0.38	SEG1
ATOM	2119	SG	CYS	138	-15.330	-4.793	1.944	1.00	0.46	SEG1
ATOM	2120	HG	CYS	138	-15.160	-4.953	2.875	1.00	0.97	SEG1
ATOM	2121	C	CYS	138	-13.819	-1.102	1.096	1.00	0.35	SEG1
ATOM	2122	O	CYS	138	-13.814	-0.249	1.964	1.00	0.40	SEG1
ATOM	2123	N	ILE	139	-12.950	-1.094	0.123	1.00	0.33	SEG1
ATOM	2124	HN	ILE	139	-12.979	-1.798	-0.557	1.00	0.33	SEG1
ATOM	2125	CA	ILE	139	-11.913	-0.019	0.049	1.00	0.35	SEG1
ATOM	2126	HA	ILE	139	-11.300	-0.030	0.938	1.00	0.37	SEG1
ATOM	2127	CB	ILE	139	-11.060	-0.363	-1.178	1.00	0.37	SEG1
ATOM	2128	HB	ILE	139	-11.686	-0.387	-2.059	1.00	0.39	SEG1
ATOM	2129	CG1	ILE	139	-10.397	-1.729	-0.980	1.00	0.37	SEG1
ATOM	2130	HG11	ILE	139	-11.136	-2.454	-0.677	1.00	0.35	SEG1
ATOM	2131	HG12	ILE	139	-9.635	-1.651	-0.218	1.00	0.40	SEG1
ATOM	2132	CG2	ILE	139	-9.968	0.694	-1.355	1.00	0.43	SEG1
ATOM	2133	HG21	ILE	139	-10.413	1.677	-1.347	1.00	1.04	SEG1
ATOM	2134	HG22	ILE	139	-9.462	0.535	-2.296	1.00	1.13	SEG1
ATOM	2135	HG23	ILE	139	-9.257	0.613	-0.546	1.00	1.14	SEG1
ATOM	2136	CD1	ILE	139	-9.757	-2.175	-2.295	1.00	0.44	SEG1
ATOM	2137	HD11	ILE	139	-8.882	-2.772	-2.088	1.00	1.04	SEG1
ATOM	2138	HD12	ILE	139	-9.473	-1.305	-2.969	1.00	1.04	SEG1
ATOM	2139	HD13	ILE	139	-10.468	-2.761	-2.859	1.00	1.12	SEG1
ATOM	2140	C	ILE	139	-12.584	1.351	-0.113	1.00	0.42	SEG1
ATOM	2141	O	ILE	139	-12.169	2.323	0.490	1.00	0.49	SEG1
ATOM	2142	N	LEU	140	-13.614	1.435	-0.918	1.00	0.44	SEG1
ATOM	2143	HN	LEU	140	-13.932	0.638	-1.393	1.00	0.41	SEG1

FIG. 2 (28 of 35)

ATOM	2144	CA	LEU	140	-14.306	2.745	-1.114	1.00	0.55	SEG1
ATOM	2145	HA	LEU	140	-13.576	3.535	-1.113	1.00	0.57	SEG1
ATOM	2146	CB	LEU	140	-14.942	2.645	-2.525	1.00	0.64	SEG1
ATOM	2147	HB1	LEU	140	-14.431	1.871	-3.076	1.00	0.65	SEG1
ATOM	2148	HB2	LEU	140	-14.805	3.585	-3.041	1.00	0.87	SEG1
ATOM	2149	CG	LEU	140	-16.445	2.313	-2.473	1.00	0.63	SEG1
ATOM	2150	HG	LEU	140	-16.635	1.638	-1.551	1.00	0.60	SEG1
ATOM	2151	CD1	LEU	140	-17.251	3.602	-2.271	1.00	0.95	SEG1
ATOM	2152	HD11	LEU	140	-16.602	4.380	-1.996	1.00	1.34	SEG1
ATOM	2153	HD12	LEU	140	-18.043	3.422	-1.558	1.00	1.62	SEG1
ATOM	2154	HD13	LEU	140	-17.679	3.915	-3.212	1.00	1.37	SEG1
ATOM	2155	CD2	LEU	140	-16.862	1.649	-3.788	1.00	0.62	SEG1
ATOM	2156	HD21	LEU	140	-17.839	1.204	-3.673	1.00	1.13	SEG1
ATOM	2157	HD22	LEU	140	-16.146	0.884	-4.048	1.00	1.26	SEG1
ATOM	2158	HD23	LEU	140	-16.895	2.391	-4.572	1.00	1.20	SEG1
ATOM	2159	C	LEU	140	-15.323	3.001	0.020	1.00	0.59	SEG1
ATOM	2160	O	LEU	140	-15.806	4.106	0.184	1.00	0.71	SEG1
ATOM	2161	N	ALA	141	-15.638	1.997	0.305	1.00	0.54	SEG1
ATOM	2162	HN	ALA	141	-15.231	1.117	0.664	1.00	0.47	SEG1
ATOM	2163	CA	ALA	141	-16.605	2.196	1.928	1.00	0.63	SEG1
ATOM	2164	HA	ALA	141	-17.502	2.676	1.571	1.00	0.73	SEG1
ATOM	2165	CB	ALA	141	-16.931	0.790	2.436	1.00	0.67	SEG1
ATOM	2166	HB1	ALA	141	-16.138	0.112	2.153	1.00	1.22	SEG1
ATOM	2167	HB2	ALA	141	-17.862	0.457	2.001	1.00	1.22	SEG1
ATOM	2168	HB3	ALA	141	-17.022	0.807	3.511	1.00	1.18	SEG1
ATOM	2169	C	ALA	141	-15.958	3.032	3.040	1.00	0.61	SEG1
ATOM	2170	O	ALA	141	-16.639	3.641	3.844	1.00	0.65	SEG1
ATOM	2171	N	GLN	142	-14.646	3.066	3.090	1.00	0.61	SEG1
ATOM	2172	HN	GLN	142	-14.118	2.569	2.433	1.00	0.64	SEG1
ATOM	2173	CA	GLN	142	-13.951	3.862	4.144	1.00	0.64	SEG1
ATOM	2174	HA	GLN	142	-14.603	4.024	4.387	1.00	0.70	SEG1
ATOM	2175	CB	GLN	142	-12.756	3.000	4.564	1.00	0.65	SEG1
ATOM	2176	HB1	GLN	142	-12.100	3.577	5.197	1.00	1.05	SEG1
ATOM	2177	HB2	GLN	142	-12.217	2.678	3.682	1.00	1.01	SEG1
ATOM	2178	CG	GLN	142	-13.256	1.772	5.332	1.00	1.11	SEG1
ATOM	2179	HG1	GLN	142	-14.028	1.279	4.761	1.00	1.54	SEG1
ATOM	2180	HG2	GLN	142	-13.658	2.085	6.285	1.00	1.56	SEG1
ATOM	2181	CD	GLN	142	-12.096	0.798	5.563	1.00	1.13	SEG1
ATOM	2182	OE1	GLN	142	-11.089	0.860	4.887	1.00	0.88	SEG1
ATOM	2183	NE2	GLN	142	-12.200	-0.110	6.496	1.00	1.49	SEG1
ATOM	2184	HE21	GLN	142	-13.014	-0.163	7.039	1.00	1.77	SEG1
ATOM	2185	HE22	GLN	142	-11.464	-0.737	6.653	1.00	1.52	SEG1
ATOM	2186	C	GLN	142	-13.480	5.202	3.566	1.00	0.65	SEG1
ATOM	2187	O	GLN	142	-12.790	5.247	2.566	1.00	0.65	SEG1
ATOM	2188	N	GLN	143	-13.855	6.292	4.190	1.00	0.74	SEG1
ATOM	2189	HN	GLN	143	-14.414	6.225	4.992	1.00	0.80	SEG1
ATOM	2190	CA	GLN	143	-13.439	7.636	3.682	1.00	0.83	SEG1
ATOM	2191	HA	GLN	143	-13.555	7.684	2.611	1.00	0.85	SEG1
ATOM	2192	CB	GLN	143	-14.388	8.631	4.353	1.00	0.97	SEG1
ATOM	2193	HB1	GLN	143	-14.037	9.637	4.178	1.00	1.46	SEG1
ATOM	2194	HB2	GLN	143	-14.416	8.439	5.416	1.00	1.09	SEG1
ATOM	2195	CG	GLN	143	-15.793	8.476	3.769	1.00	1.27	SEG1
ATOM	2196	HG1	GLN	143	-16.159	7.480	3.968	1.00	1.58	SEG1
ATOM	2197	HG2	GLN	143	-15.760	8.642	2.701	1.00	1.81	SEG1
ATOM	2198	CD	GLN	143	-16.727	9.499	4.417	1.00	1.60	SEG1
ATOM	2199	OE1	GLN	143	-16.864	9.534	5.624	1.00	1.78	SEG1
ATOM	2200	NE2	GLN	143	-17.377	10.343	3.663	1.00	2.32	SEG1
ATOM	2201	HE21	GLN	143	-17.265	10.317	2.689	1.00	2.64	SEG1
ATOM	2202	HE22	GLN	143	-17.978	11.002	4.070	1.00	2.75	SEG1
ATOM	2203	C	GLN	143	-11.984	7.936	4.075	1.00	0.85	SEG1
ATOM	2204	O	GLN	143	-11.506	7.443	5.078	1.00	0.92	SEG1
ATOM	2205	N	PRO	144	-11.317	8.740	3.272	1.00	0.88	SEG1
ATOM	2206	CA	PRO	144	-9.905	9.093	3.561	1.00	0.95	SEG1
ATOM	2207	HA	PRO	144	-9.322	8.206	3.736	1.00	1.00	SEG1
ATOM	2208	CB	PRO	144	-9.435	9.784	2.283	1.00	1.01	SEG1
ATOM	2209	HB1	PRO	144	-8.960	9.074	1.624	1.00	1.08	SEG1
ATOM	2210	HB2	PRO	144	-8.757	10.592	2.522	1.00	1.06	SEG1
ATOM	2211	CG	PRO	144	-10.680	10.315	1.651	1.00	1.03	SEG1
ATOM	2212	HG1	PRO	144	-10.574	10.327	0.577	1.00	1.13	SEG1
ATOM	2213	HG2	PRO	144	-10.882	11.313	2.015	1.00	1.06	SEG1
ATOM	2214	CD	PRO	144	-11.798	9.385	2.039	1.00	0.96	SEG1
ATOM	2215	HD2	PRO	144	-12.705	9.943	2.228	1.00	1.00	SEG1
ATOM	2216	HD1	PRO	144	-11.958	8.644	1.273	1.00	1.01	SEG1
ATOM	2217	C	PRO	144	-9.821	10.047	4.759	1.00	0.99	SEG1
ATOM	2218	O	PRO	144	-10.815	10.594	5.199	1.00	1.45	SEG1
ATOM	2219	N	ASP	145	-8.638	10.250	5.283	1.00	0.92	SEG1
ATOM	2220	HN	ASP	145	-7.856	9.800	4.904	1.00	1.14	SEG1

FIG. 2 (29 of 35)

ATOM	2221	CA	ASP	145	-8.471	11.168	6.448	1.00	1.02	SEG1
ATOM	2222	HA	ASP	145	-9.243	11.921	6.450	1.00	1.13	SEG1
ATOM	2223	CB	ASP	145	-8.605	10.277	7.683	1.00	1.23	SEG1
ATOM	2224	HB1	ASP	145	-7.754	9.617	7.744	1.00	1.40	SEG1
ATOM	2225	HB2	ASP	145	-9.511	9.693	7.607	1.00	1.68	SEG1
ATOM	2226	CG	ASP	145	-8.663	11.150	8.938	1.00	1.86	SEG1
ATOM	2227	OD1	ASP	145	-9.712	11.193	9.559	1.00	2.54	SEG1
ATOM	2228	OD2	ASP	145	-7.656	11.761	9.257	1.00	2.40	SEG1
ATOM	2229	C	ASP	145	-7.086	11.818	6.405	1.00	0.94	SEG1
ATOM	2230	O	ASP	145	-6.127	11.296	6.943	1.00	0.94	SEG1
ATOM	2231	N	APG	146	-6.980	12.951	5.763	1.00	1.05	SEG1
ATOM	2232	HN	APG	146	-7.770	13.341	5.337	1.00	1.19	SEG1
ATOM	2233	CA	APG	146	-5.662	13.650	5.669	1.00	1.10	SEG1
ATOM	2234	HA	APG	146	-4.856	12.934	5.648	1.00	1.16	SEG1
ATOM	2235	CB	APG	146	-5.720	14.400	4.339	1.00	1.42	SEG1
ATOM	2236	HB1	ARG	146	-4.879	15.074	4.269	1.00	1.92	SEG1
ATOM	2237	HB2	ARG	146	-6.639	14.965	4.287	1.00	1.98	SEG1
ATOM	2238	CG	APG	146	-5.668	13.403	3.181	1.00	1.52	SEG1
ATOM	2239	HG1	APG	146	-6.469	12.687	3.283	1.00	1.96	SEG1
ATOM	2240	HG2	ARG	146	-4.719	12.886	3.193	1.00	1.91	SEG1
ATOM	2241	CD	APG	146	-5.826	14.157	1.858	1.00	1.77	SEG1
ATOM	2242	HD1	ARG	146	-6.739	14.730	1.864	1.00	2.23	SEG1
ATOM	2243	HD2	APG	146	-5.818	13.463	1.028	1.00	1.99	SEG1
ATOM	2244	NE	ARG	146	-4.645	15.074	1.789	1.00	2.23	SEG1
ATOM	2245	HE	APG	146	-3.845	14.864	2.315	1.00	2.67	SEG1
ATOM	2246	CZ	ARG	146	-4.656	16.158	1.037	1.00	2.70	SEG1
ATOM	2247	NH1	ARG	146	-3.598	16.921	1.010	1.00	3.55	SEG1
ATOM	2248	HH11	ARG	146	-2.794	16.682	1.556	1.00	3.97	SEG1
ATOM	2249	HH12	ARG	146	-3.590	17.746	0.443	1.00	3.99	SEG1
ATOM	2250	NH2	ARG	146	-5.703	16.487	0.313	1.00	2.82	SEG1
ATOM	2251	HH21	ARG	146	-6.522	15.917	0.317	1.00	2.74	SEG1
ATOM	2252	HH22	ARG	146	-5.679	17.317	-0.245	1.00	3.37	SEG1
ATOM	2253	C	ARG	146	-5.469	14.641	6.829	1.00	0.99	SEG1
ATOM	2254	O	ARG	146	-4.407	15.215	6.981	1.00	0.96	SEG1
ATOM	2255	N	LEU	147	-6.480	14.856	7.640	1.00	1.03	SEG1
ATOM	2256	HN	LEU	147	-7.330	14.391	7.500	1.00	1.10	SEG1
ATOM	2257	CA	LEU	147	-6.336	15.821	8.777	1.00	1.07	SEG1
ATOM	2258	HA	LEU	147	-6.140	16.813	8.406	1.00	1.17	SEG1
ATOM	2259	CB	LEU	147	-7.683	15.794	9.504	1.00	1.28	SEG1
ATOM	2260	HB1	LEU	147	-7.603	16.343	10.431	1.00	1.65	SEG1
ATOM	2261	HB2	LEU	147	-7.958	14.770	9.716	1.00	1.48	SEG1
ATOM	2262	CG	LEU	147	-8.755	16.439	8.625	1.00	1.81	SEG1
ATOM	2263	HG	LEU	147	-8.746	15.975	7.649	1.00	2.37	SEG1
ATOM	2264	CD1	LEU	147	-10.128	16.239	9.268	1.00	2.27	SEG1
ATOM	2265	HD11	LEU	147	-10.862	16.833	8.743	1.00	2.67	SEG1
ATOM	2266	HD12	LEU	147	-10.089	16.548	10.302	1.00	2.60	SEG1
ATOM	2267	HD13	LEU	147	-10.403	15.197	9.214	1.00	2.67	SEG1
ATOM	2268	CD2	LEU	147	-8.471	17.936	8.485	1.00	2.31	SEG1
ATOM	2269	HD21	LEU	147	-9.400	18.467	8.341	1.00	2.50	SEG1
ATOM	2270	HD22	LEU	147	-7.827	18.103	7.634	1.00	2.84	SEG1
ATOM	2271	HD23	LEU	147	-7.985	18.296	9.380	1.00	2.68	SEG1
ATOM	2272	C	LEU	147	-5.217	15.372	9.720	1.00	0.95	SEG1
ATOM	2273	O	LEU	147	-4.415	16.168	10.171	1.00	1.02	SEG1
ATOM	2274	N	ARG	148	-5.166	14.103	10.020	1.00	0.90	SEG1
ATOM	2275	HN	ARG	148	-5.829	13.489	9.643	1.00	0.90	SEG1
ATOM	2276	CA	ARG	148	-4.108	13.582	10.939	1.00	1.02	SEG1
ATOM	2277	HA	ARG	148	-3.994	14.241	11.785	1.00	1.19	SEG1
ATOM	2278	CB	ARG	148	-4.616	12.221	11.414	1.00	1.08	SEG1
ATOM	2279	HB1	ARG	148	-3.828	11.707	11.943	1.00	1.58	SEG1
ATOM	2280	HB2	ARG	148	-4.920	11.632	10.560	1.00	1.16	SEG1
ATOM	2281	CG	ARG	148	-5.809	12.416	12.349	1.00	1.69	SEG1
ATOM	2282	HG1	ARG	148	-6.602	12.921	11.818	1.00	2.05	SEG1
ATOM	2283	HG2	ARG	148	-5.507	13.011	13.198	1.00	2.34	SEG1
ATOM	2284	CD	ARG	148	-6.309	11.052	12.827	1.00	1.94	SEG1
ATOM	2285	HD1	ARG	148	-5.530	10.532	13.364	1.00	2.37	SEG1
ATOM	2286	HD2	ARG	148	-6.650	10.461	11.987	1.00	2.25	SEG1
ATOM	2287	NE	ARG	148	-7.443	11.363	13.746	1.00	2.31	SEG1
ATOM	2288	HE	ARG	148	-8.363	11.204	13.446	1.00	2.69	SEG1
ATOM	2289	CZ	ARG	148	-7.232	11.843	14.957	1.00	2.75	SEG1
ATOM	2290	NH1	ARG	148	-8.253	12.092	15.731	1.00	3.51	SEG1
ATOM	2291	HH11	ARG	148	-9.181	11.918	15.404	1.00	3.86	SEG1
ATOM	2292	HH12	ARG	148	-8.106	12.458	16.650	1.00	3.96	SEG1
ATOM	2293	NH2	ARG	148	-6.018	12.081	25.402	1.00	2.95	SEG1
ATOM	2294	HH21	ARG	148	-5.222	11.899	14.825	1.00	2.88	SEG1
ATOM	2295	HH22	ARG	148	-5.890	12.446	16.324	1.00	3.50	SEG1
ATOM	2296	C	ARG	148	-2.769	13.418	10.215	1.00	1.06	SEG1
ATOM	2297	O	ARG	148	-1.730	13.342	10.844	1.00	1.28	SEG1

FIG. 2 (30 of 35)

ATOM	2298	N	ASP	149	-2.782	13	330	8.904	1.00	1.01	SEG1
ATOM	2299	HN	ASP	149	-3.631	13	370	8.423	1.00	0.94	SEG1
ATOM	2300	CA	ASP	149	-1.501	13	135	8.164	1.00	1.21	SEG1
ATOM	2301	HA	ASP	149	-0.688	12	999	8.853	1.00	1.42	SEG1
ATOM	2302	CB	ASP	149	-1.708	11	849	7.361	1.00	1.48	SEG1
ATOM	2303	HB1	ASP	149	-0.862	11	692	6.711	1.00	1.86	SEG1
ATOM	2304	HB2	ASP	149	-2.606	11	938	6.767	1.00	1.80	SEG1
ATOM	2305	CG	ASP	149	-1.843	10	655	8.316	1.00	1.62	SEG1
ATOM	2306	OD1	ASP	149	-2.324	9	624	7.874	1.00	2.12	SEG1
ATOM	2307	OD2	ASP	149	-1.471	10	790	9.473	1.00	2.10	SEG1
ATOM	2308	C	ASP	149	-1.205	14	314	7.233	1.00	1.03	SEG1
ATOM	2309	O	ASP	149	-0.684	14	133	6.147	1.00	0.90	SEG1
ATOM	2310	N	GLU	150	-1.521	15	516	7.646	1.00	1.07	SEG1
ATOM	2311	HN	GLU	150	-1.934	15	638	8.527	1.00	1.23	SEG1
ATOM	2312	CA	GLU	150	-1.242	16	703	6.776	1.00	0.95	SEG1
ATOM	2313	HA	GLU	150	-1.797	16	629	5.855	1.00	0.95	SEG1
ATOM	2314	CB	GLU	150	-1.718	17	917	7.574	1.00	1.11	SEG1
ATOM	2315	HB1	GLU	150	-1.378	18	821	7.092	1.00	1.44	SEG1
ATOM	2316	HB2	GLU	150	-1.315	17	869	8.576	1.00	1.75	SEG1
ATOM	2317	CG	GLU	150	-3.246	17	919	7.638	1.00	1.35	SEG1
ATOM	2318	HG1	GLU	150	-3.589	17	017	8.120	1.00	1.98	SEG1
ATOM	2319	HG2	GLU	150	-3.646	17	968	6.635	1.00	1.71	SEG1
ATOM	2320	CD	GLU	150	-3.720	19	135	8.436	1.00	1.41	SEG1
ATOM	2321	OE1	GLU	150	-3.787	19	033	9.650	1.00	1.66	SEG1
ATOM	2322	OE2	GLU	150	-4.008	20	148	7.819	1.00	1.91	SEG1
ATOM	2323	C	GLU	150	0.260	16	808	6.485	1.00	0.77	SEG1
ATOM	2324	O	GLU	150	0.661	17	120	5.380	1.00	0.66	SEG1
ATOM	2325	N	GLU	151	1.092	16	546	7.466	1.00	0.79	SEG1
ATOM	2326	HN	GLU	151	0.744	16	293	8.347	1.00	0.90	SEG1
ATOM	2327	CA	GLU	151	2.571	16	627	7.239	1.00	0.70	SEG1
ATOM	2328	HA	GLU	151	2.853	17	630	6.967	1.00	0.73	SEG1
ATOM	2329	CB	GLU	151	3.216	16	244	8.573	1.00	0.82	SEG1
ATOM	2330	HB1	GLU	151	4.279	16	114	8.434	1.00	1.27	SEG1
ATOM	2331	HB2	GLU	151	2.783	15	321	8.931	1.00	1.15	SEG1
ATOM	2332	CG	GLU	151	2.971	17	356	9.600	1.00	1.41	SEG1
ATOM	2333	HG1	GLU	151	1.909	17	490	9.736	1.00	1.98	SEG1
ATOM	2334	HG2	GLU	151	3.404	18	277	9.239	1.00	1.90	SEG1
ATOM	2335	CD	GLU	151	3.613	16	988	10.948	1.00	1.66	SEG1
ATOM	2336	OE1	GLU	151	3.401	17	725	11.898	1.00	2.04	SEG1
ATOM	2337	OE2	GLU	151	4.305	15	982	11.012	1.00	2.22	SEG1
ATOM	2338	C	GLU	151	2.990	15	643	6.144	1.00	0.54	SEG1
ATOM	2339	O	GLU	151	3.816	15	950	5.305	1.00	0.48	SEG1
ATOM	2340	N	LEU	152	2.415	14	468	6.142	1.00	0.54	SEG1
ATOM	2341	HN	LEU	152	1.747	14	252	6.825	1.00	0.63	SEG1
ATOM	2342	CA	LEU	152	2.764	13	460	5.097	1.00	0.46	SEG1
ATOM	2343	HA	LEU	152	3.830	13	295	5.078	1.00	0.44	SEG1
ATOM	2344	CB	LEU	152	2.046	12	171	5.499	1.00	0.55	SEG1
ATOM	2345	HB1	LEU	152	2.098	11	463	4.686	1.00	0.56	SEG1
ATOM	2346	HB2	LEU	152	1.011	12	390	5.717	1.00	0.63	SEG1
ATOM	2347	CG	LEU	152	2.710	11	569	6.736	1.00	0.59	SEG1
ATOM	2348	HG	LEU	152	2.753	12	310	7.522	1.00	0.62	SEG1
ATOM	2349	CD1	LEU	152	1.897	10	365	7.209	1.00	0.70	SEG1
ATOM	2350	HD11	LEU	152	0.850	10	538	7.013	1.00	1.18	SEG1
ATOM	2351	HD12	LEU	152	2.046	10	223	8.269	1.00	1.17	SEG1
ATOM	2352	HD13	LEU	152	2.222	9	481	6.677	1.00	1.38	SEG1
ATOM	2353	CD2	LEU	152	4.126	11	112	6.379	1.00	0.56	SEG1
ATOM	2354	HD21	LEU	152	4.492	10	443	7.144	1.00	1.07	SEG1
ATOM	2355	HD22	LEU	152	4.777	11	971	6.311	1.00	1.17	SEG1
ATOM	2356	HD23	LEU	152	4.108	10	596	5.430	1.00	1.20	SEG1
ATOM	2357	C	LEU	152	2.278	13	931	3.728	1.00	0.46	SEG1
ATOM	2358	O	LEU	152	2.925	13	708	2.725	1.00	0.46	SEG1
ATOM	2359	N	ALA	153	1.138	14	577	3.682	1.00	0.54	SEG1
ATOM	2360	HN	ALA	153	0.636	14	739	4.509	1.00	0.57	SEG1
ATOM	2361	CA	ALA	153	0.599	15	060	2.371	1.00	0.61	SEG1
ATOM	2362	HA	ALA	153	0.408	14	225	1.715	1.00	0.65	SEG1
ATOM	2363	CB	ALA	153	-0.713	15	771	2.711	1.00	0.71	SEG1
ATOM	2364	HB1	ALA	153	-1.536	15	081	2.600	1.00	1.20	SEG1
ATOM	2365	HB2	ALA	153	-0.851	16	609	2.043	1.00	1.25	SEG1
ATOM	2366	HB3	ALA	153	-0.676	16	126	3.730	1.00	1.26	SEG1
ATOM	2367	C	ALA	153	1.587	16	033	1.723	1.00	0.58	SEG1
ATOM	2368	O	ALA	153	1.810	15	997	0.527	1.00	0.63	SEG1
ATOM	2369	N	GLU	154	2.191	16	893	2.502	1.00	0.53	SEG1
ATOM	2370	HN	GLU	154	2.005	16	899	3.464	1.00	0.52	SEG1
ATOM	2371	CA	GLU	154	3.183	17	844	1.924	1.00	0.55	SEG1
ATOM	2372	HA	GLU	154	2.764	18	359	1.073	1.00	0.62	SEG1
ATOM	2373	CB	GLU	154	3.501	18	837	3.044	1.00	0.56	SEG1
ATOM	2374	HB1	GLU	154	4.339	19	451	2.752	1.00	1.06	SEG1

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ATOM	2375	HB1	GLU	154	3.748	18.294	3.945	1.00	0.99	SEG1
ATOM	2376	CG	GLU	154	2.283	19.729	3.300	1.00	1.00	SEG1
ATOM	2377	HG1	GLU	154	1.446	19.112	3.592	1.00	1.52	SEG1
ATOM	2378	HG2	GLU	154	2.034	20.265	2.396	1.00	1.58	SEG1
ATOM	2379	CD	GLU	154	2.583	20.733	4.423	1.00	1.17	SEG1
ATOM	2380	OE1	GLU	154	1.777	21.631	4.612	1.00	1.60	SEG1
ATOM	2381	OE2	GLU	154	3.606	20.591	5.077	1.00	1.77	SEG1
ATOM	2382	C	GLU	154	4.440	17.073	1.518	1.00	0.51	SEG1
ATOM	2383	O	GLU	154	5.039	17.330	0.491	1.00	0.55	SEG1
ATOM	2384	N	LEU	155	4.839	16.127	2.332	1.00	0.45	SEG1
ATOM	2385	HN	LEU	155	4.332	15.950	3.153	1.00	0.44	SEG1
ATOM	2386	CA	LEU	155	6.059	15.323	2.021	1.00	0.43	SEG1
ATOM	2387	HA	LEU	155	6.921	15.964	1.958	1.00	0.47	SEG1
ATOM	2388	CB	LEU	155	6.213	14.356	3.202	1.00	0.40	SEG1
ATOM	2389	HB1	LEU	155	5.335	13.731	3.270	1.00	0.40	SEG1
ATOM	2390	HB2	LEU	155	6.324	14.921	4.116	1.00	0.43	SEG1
ATOM	2391	CG	LEU	155	7.447	13.471	2.994	1.00	0.42	SEG1
ATOM	2392	HG	LEU	155	8.195	14.025	2.445	1.00	0.51	SEG1
ATOM	2393	CD1	LEU	155	8.016	13.060	4.353	1.00	0.42	SEG1
ATOM	2394	HD11	LEU	155	7.475	12.202	4.724	1.00	1.09	SEG1
ATOM	2395	HD12	LEU	155	7.914	13.879	5.050	1.00	1.01	SEG1
ATOM	2396	HD13	LEU	155	9.060	12.808	4.244	1.00	1.13	SEG1
ATOM	2397	CD2	LEU	155	7.054	12.221	2.208	1.00	0.51	SEG1
ATOM	2398	HD21	LEU	155	6.036	11.950	2.447	1.00	1.03	SEG1
ATOM	2399	HD22	LEU	155	7.714	11.408	2.472	1.00	1.18	SEG1
ATOM	2400	HD23	LEU	155	7.134	12.420	1.150	1.00	1.19	SEG1
ATOM	2401	C	LEU	155	5.875	14.554	0.709	1.00	0.46	SEG1
ATOM	2402	O	LEU	155	6.720	14.597	-0.165	1.00	0.51	SEG1
ATOM	2403	N	GLU	156	4.777	13.851	0.568	1.00	0.48	SEG1
ATOM	2404	HN	GLU	156	4.115	13.832	1.289	1.00	0.47	SEG1
ATOM	2405	CA	GLU	156	4.543	13.076	-0.689	1.00	0.55	SEG1
ATOM	2406	HA	GLU	156	5.339	12.362	-0.829	1.00	0.54	SEG1
ATOM	2407	CB	GLU	156	3.215	12.329	-0.491	1.00	0.60	SEG1
ATOM	2408	HB1	GLU	156	3.292	11.687	0.374	1.00	0.58	SEG1
ATOM	2409	HB2	GLU	156	3.011	11.728	-1.365	1.00	0.68	SEG1
ATOM	2410	CG	GLU	156	2.070	13.318	-0.281	1.00	0.64	SEG1
ATOM	2411	HG1	GLU	156	2.015	13.994	-1.120	1.00	0.72	SEG1
ATOM	2412	HG2	GLU	156	2.245	13.874	0.623	1.00	0.57	SEG1
ATOM	2413	CD	GLU	156	0.751	12.553	-0.159	1.00	0.74	SEG1
ATOM	2414	OE1	GLU	156	0.015	12.523	-1.132	1.00	1.30	SEG1
ATOM	2415	OE2	GLU	156	0.500	12.011	0.904	1.00	1.37	SEG1
ATOM	2416	C	GLU	156	4.478	14.025	-1.889	1.00	0.63	SEG1
ATOM	2417	O	GLU	156	4.938	13.700	-2.968	1.00	0.67	SEG1
ATOM	2418	N	ASP	157	3.922	15.199	-1.707	1.00	0.66	SEG1
ATOM	2419	HN	ASP	157	3.566	15.441	-0.826	1.00	0.63	SEG1
ATOM	2420	CA	ASP	157	3.842	16.169	-2.842	1.00	0.76	SEG1
ATOM	2421	HA	ASP	157	3.408	15.697	-3.708	1.00	0.81	SEG1
ATOM	2422	CB	ASP	157	2.940	17.304	-2.351	1.00	0.81	SEG1
ATOM	2423	HB1	ASP	157	2.980	18.123	-3.053	1.00	0.88	SEG1
ATOM	2424	HB2	ASP	157	3.283	17.644	-1.384	1.00	0.76	SEG1
ATOM	2425	CG	ASP	157	1.496	16.805	-2.235	1.00	0.84	SEG1
ATOM	2426	OD1	ASP	157	1.133	15.914	-2.985	1.00	1.26	SEG1
ATOM	2427	OD2	ASP	157	0.778	17.324	-1.397	1.00	1.49	SEG1
ATOM	2428	C	ASP	157	5.237	16.691	-3.171	1.00	0.76	SEG1
ATOM	2429	O	ASP	157	5.597	16.841	-4.323	1.00	0.82	SEG1
ATOM	2430	N	ALA	158	6.029	16.950	-2.165	1.00	0.70	SEG1
ATOM	2431	HN	ALA	158	5.714	16.806	-1.247	1.00	0.66	SEG1
ATOM	2432	CA	ALA	158	7.414	17.444	-2.412	1.00	0.74	SEG1
ATOM	2433	HA	ALA	158	7.396	18.312	-3.051	1.00	0.82	SEG1
ATOM	2434	CB	ALA	158	7.969	17.814	-1.034	1.00	0.72	SEG1
ATOM	2435	HB1	ALA	158	7.160	18.133	-0.394	1.00	1.34	SEG1
ATOM	2436	HB2	ALA	158	8.684	18.617	-1.138	1.00	1.21	SEG1
ATOM	2437	HB3	ALA	158	8.455	16.954	-0.599	1.00	1.17	SEG1
ATOM	2438	C	ALA	158	8.240	16.325	-3.050	1.00	0.70	SEG1
ATOM	2439	O	ALA	158	9.052	16.559	-3.926	1.00	0.77	SEG1
ATOM	2440	N	LEU	159	8.031	15.109	-2.612	1.00	0.62	SEG1
ATOM	2441	HN	LEU	159	7.369	14.957	-1.905	1.00	0.58	SEG1
ATOM	2442	CA	LEU	159	8.797	13.956	-3.181	1.00	0.60	SEG1
ATOM	2443	HA	LEU	159	9.850	14.072	-2.989	1.00	0.63	SEG1
ATOM	2444	CB	LEU	159	8.265	12.717	-2.448	1.00	0.53	SEG1
ATOM	2445	HB1	LEU	159	7.206	12.617	-2.633	1.00	0.53	SEG1
ATOM	2446	HB2	LEU	159	8.437	12.825	-1.388	1.00	0.51	SEG1
ATOM	2447	CG	LEU	159	8.988	11.465	-2.956	1.00	0.54	SEG1
ATOM	2448	HG	LEU	159	8.963	11.453	-4.037	1.00	0.65	SEG1
ATOM	2449	CD1	LEU	159	10.446	11.479	-2.482	1.00	0.76	SEG1
ATOM	2450	HD11	LEU	159	10.731	12.485	-2.214	1.00	1.37	SEG1
ATOM	2451	HD12	LEU	159	11.086	11.124	-3.276	1.00	1.37	SEG1

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ATOM	2451	HD13	LEU	159	10.552	10.835	-1.621	1.00	1.16	SEG1
ATOM	2453	CD2	LEU	159	8.283	10.221	-2.416	1.00	0.58	SEG1
ATOM	2454	HD21	LEU	159	8.485	10.124	-1.360	1.00	1.22	SEG1
ATOM	2455	HD22	LEU	159	8.647	9.347	-2.935	1.00	1.13	SEG1
ATOM	2456	HD23	LEU	159	7.218	10.315	-2.572	1.00	1.21	SEG1
ATOM	2457	C	LEU	159	8.534	13.836	-4.688	1.00	0.67	SEG1
ATOM	2458	O	LEU	159	9.438	13.579	-5.460	1.00	0.71	SEG1
ATOM	2459	N	AFG	160	7.303	14.018	-5.111	1.00	0.70	SEG1
ATOM	2460	HN	AFG	160	6.589	14.224	-4.467	1.00	0.68	SEG1
ATOM	2461	CA	AFG	160	6.988	13.912	-6.572	1.00	0.80	SEG1
ATOM	2462	HA	AFG	160	7.205	12.921	-6.932	1.00	0.80	SEG1
ATOM	2463	CB	AFG	160	5.486	14.197	-6.687	1.00	0.86	SEG1
ATOM	2464	HB1	AFG	160	5.277	15.186	-6.310	1.00	1.08	SEG1
ATOM	2465	HB2	AFG	160	4.938	13.467	-6.108	1.00	1.05	SEG1
ATOM	2466	CG	AFG	160	5.054	14.112	-8.156	1.00	1.22	SEG1
ATOM	2467	HG1	AFG	160	5.258	13.121	-8.533	1.00	1.67	SEG1
ATOM	2468	HG2	AFG	160	5.604	14.839	-8.736	1.00	1.51	SEG1
ATOM	2469	CD	AFG	160	3.552	14.396	-8.264	1.00	1.41	SEG1
ATOM	2470	HD1	ARG	160	3.334	15.392	-7.911	1.00	1.63	SEG1
ATOM	2471	HD2	ARG	160	2.992	13.665	-7.699	1.00	1.87	SEG1
ATOM	2472	NE	AFG	160	3.230	14.290	-9.723	1.00	1.88	SEG1
ATOM	2473	HE	AFG	160	3.876	13.972	-10.330	1.00	2.40	SEG1
ATOM	2474	CZ	AFG	160	2.091	14.750	-10.208	1.00	2.25	SEG1
ATOM	2475	NH1	AFG	160	1.851	14.638	-11.487	1.00	3.04	SEG1
ATOM	2476	NH11	AFG	160	2.525	14.208	-12.087	1.00	3.44	SEG1
ATOM	2477	NH12	ARG	160	0.992	14.983	-11.865	1.00	3.46	SEG1
ATOM	2478	NH2	ARG	160	1.193	15.318	-9.436	1.00	2.40	SEG1
ATOM	2479	NH21	AFG	160	1.357	15.414	-8.455	1.00	2.30	SEG1
ATOM	2480	NH22	ARG	160	0.340	15.656	-9.833	1.00	2.98	SEG1
ATOM	2481	C	ARG	160	7.800	14.953	-7.354	1.00	0.90	SEG1
ATOM	2482	O	ARG	160	8.332	14.673	-8.412	1.00	0.97	SEG1
ATOM	2483	N	ASN	161	7.904	16.145	-6.827	1.00	0.92	SEG1
ATOM	2484	HN	ASN	161	7.470	16.332	-5.969	1.00	0.87	SEG1
ATOM	2485	CA	ASN	161	8.690	17.214	-7.516	1.00	1.04	SEG1
ATOM	2486	HA	ASN	161	8.312	17.378	-8.511	1.00	1.11	SEG1
ATOM	2487	CB	ASN	161	8.495	18.472	-6.666	1.00	1.08	SEG1
ATOM	2488	HB1	ASN	161	9.172	19.244	-7.002	1.00	1.35	SEG1
ATOM	2489	HB2	ASN	161	8.697	18.241	-5.631	1.00	1.08	SEG1
ATOM	2490	CG	ASN	161	7.053	18.966	-6.806	1.00	1.59	SEG1
ATOM	2491	OD1	ASN	161	6.414	18.733	-7.812	1.00	2.33	SEG1
ATOM	2492	ND2	ASN	161	6.510	19.642	-5.831	1.00	1.73	SEG1
ATOM	2493	HD21	ASN	161	7.024	19.827	-5.017	1.00	1.70	SEG1
ATOM	2494	HD22	ASN	161	5.588	19.967	-5.913	1.00	2.23	SEG1
ATOM	2495	C	ASN	161	10.169	16.820	-7.563	1.00	1.05	SEG1
ATOM	2496	O	ASN	161	10.850	17.034	-8.548	1.00	1.13	SEG1
ATOM	2497	N	LEU	162	10.663	16.249	-6.497	1.00	0.97	SEG1
ATOM	2498	HN	LEU	162	10.083	16.096	-5.722	1.00	0.90	SEG1
ATOM	2499	CA	LEU	162	12.099	15.831	-6.451	1.00	1.01	SEG1
ATOM	2500	HA	LEU	162	12.742	16.692	-6.547	1.00	1.10	SEG1
ATOM	2501	CB	LEU	162	12.285	15.190	-5.070	1.00	0.95	SEG1
ATOM	2502	HB1	LEU	162	11.617	14.348	-4.973	1.00	1.27	SEG1
ATOM	2503	HB2	LEU	162	12.059	15.918	-4.305	1.00	1.20	SEG1
ATOM	2504	CG	LEU	162	13.732	14.713	-4.908	1.00	1.50	SEG1
ATOM	2505	HG	LEU	162	13.999	14.089	-5.749	1.00	2.22	SEG1
ATOM	2506	CD1	LEU	162	14.672	15.917	-4.852	1.00	1.92	SEG1
ATOM	2507	HD11	LEU	162	15.695	15.573	-4.813	1.00	2.34	SEG1
ATOM	2508	HD12	LEU	162	14.454	16.503	-3.971	1.00	2.30	SEG1
ATOM	2509	HD13	LEU	162	14.531	16.525	-5.732	1.00	2.35	SEG1
ATOM	2510	CD2	LEU	162	13.861	13.906	-3.613	1.00	1.89	SEG1
ATOM	2511	HD21	LEU	162	13.042	13.207	-3.541	1.00	2.44	SEG1
ATOM	2512	HD22	LEU	162	13.837	14.577	-2.766	1.00	2.30	SEG1
ATOM	2513	HD23	LEU	162	14.796	13.366	-3.618	1.00	2.12	SEG1
ATOM	2514	C	LEU	162	12.398	14.813	-7.555	1.00	1.00	SEG1
ATOM	2515	O	LEU	162	13.418	14.886	-8.215	1.00	1.08	SEG1
ATOM	2516	N	LYS	163	11.522	13.861	-7.751	1.00	0.94	SEG1
ATOM	2517	HN	LYS	163	10.711	13.821	-7.201	1.00	0.89	SEG1
ATOM	2518	CA	LYS	163	11.759	12.833	-8.808	1.00	0.96	SEG1
ATOM	2519	HA	LYS	163	12.800	12.550	-8.828	1.00	0.98	SEG1
ATOM	2520	CB	LYS	163	10.900	11.630	-8.406	1.00	0.95	SEG1
ATOM	2521	HB1	LYS	163	9.858	11.914	-8.413	1.00	1.38	SEG1
ATOM	2522	HB2	LYS	163	11.177	11.301	-7.414	1.00	1.07	SEG1
ATOM	2523	CG	LYS	163	11.120	10.491	-9.408	1.00	1.47	SEG1
ATOM	2524	HG1	LYS	163	12.163	10.211	-9.416	1.00	1.90	SEG1
ATOM	2525	HG2	LYS	163	10.830	10.822	-10.395	1.00	2.04	SEG1
ATOM	2526	CD	LYS	163	10.271	9.282	-9.607	1.00	1.66	SEG1
ATOM	2527	HD1	LYS	163	9.224	9.543	-9.061	1.00	1.94	SEG1
ATOM	2528	HD2	LYS	163	10.519	8.991	-7.997	1.00	1.60	SEG1

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ATOM	2529	CE	LYS	163	10.552	8.113	-9.956	1.00	2.53	SEG1
ATOM	2530	HE1	LYS	163	11.609	8.036	-10.158	1.00	2.77	SEG1
ATOM	2531	HE2	LYS	163	9.998	8.236	-10.878	1.00	3.08	SEG1
ATOM	2532	NZ	LYS	163	10.082	6.907	-9.219	1.00	3.06	SEG1
ATOM	2533	H21	LYS	163	9.068	7.001	-9.211	1.00	3.11	SEG1
ATOM	2534	H22	LYS	163	10.613	6.819	-8.328	1.00	2.52	SEG1
ATOM	2535	H23	LYS	163	10.237	6.061	-9.802	1.00	3.54	SEG1
ATOM	2536	C	LYS	163	11.325	13.370	-10.172	1.00	1.08	SEG1
ATOM	2537	O	LYS	163	10.223	13.858	-10.332	1.00	1.51	SEG1
ATOM	2538	N	CYS	164	12.185	13.278	-11.154	1.00	1.37	SEG1
ATOM	2539	HN	CYS	164	13.065	12.878	-10.996	1.00	1.81	SEG1
ATOM	2540	CA	CYS	164	11.828	13.776	-12.516	1.00	1.52	SEG1
ATOM	2541	HA	CYS	164	11.144	14.606	-12.448	1.00	1.31	SEG1
ATOM	2542	CB	CYS	164	13.149	14.234	-13.135	1.00	2.17	SEG1
ATOM	2543	HB1	CYS	164	13.791	13.377	-13.285	1.00	2.58	SEG1
ATOM	2544	HB2	CYS	164	13.634	14.936	-12.472	1.00	2.61	SEG1
ATOM	2545	SG	CYS	164	12.827	15.032	-14.727	1.00	2.71	SEG1
ATOM	2546	HG	CYS	164	12.479	15.910	-14.555	1.00	3.08	SEG1
ATOM	2547	C	CYS	164	11.217	12.641	-13.340	1.00	2.18	SEG1
ATOM	2548	O	CYS	164	11.906	11.728	-13.755	1.00	2.81	SEG1
ATOM	2549	N	GLY	165	9.930	12.688	-13.573	1.00	2.69	SEG1
ATOM	2550	HN	GLY	165	9.397	13.432	-13.222	1.00	2.77	SEG1
ATOM	2551	CA	GLY	165	9.269	11.608	-14.364	1.00	3.67	SEG1
ATOM	2552	HA1	GLY	165	8.369	11.292	-13.859	1.00	4.30	SEG1
ATOM	2553	HA2	GLY	165	9.944	10.768	-14.450	1.00	4.00	SEG1
ATOM	2554	C	GLY	165	8.915	12.119	-15.765	1.00	3.89	SEG1
ATOM	2555	O	GLY	165	7.759	12.181	-16.140	1.00	3.93	SEG1
ATOM	2556	N	SER	166	9.909	12.471	-16.543	1.00	4.51	SEG1
ATOM	2557	HN	SER	166	10.828	12.400	-16.216	1.00	4.82	SEG1
ATOM	2558	CA	SER	166	9.654	12.967	-17.933	1.00	5.11	SEG1
ATOM	2559	HA	SER	166	8.668	13.401	-18.006	1.00	5.04	SEG1
ATOM	2560	CB	SER	166	10.715	14.044	-18.176	1.00	5.85	SEG1
ATOM	2561	HB1	SER	166	11.184	13.880	-19.140	1.00	6.15	SEG1
ATOM	2562	HB2	SER	166	11.465	13.997	-17.405	1.00	6.27	SEG1
ATOM	2563	OG	SER	166	10.095	15.323	-18.149	1.00	6.09	SEG1
ATOM	2564	HG	SER	166	10.696	15.935	-17.717	1.00	6.52	SEG1
ATOM	2565	C	SER	166	9.818	11.829	-18.953	1.00	5.69	SEG1
ATOM	2566	O	SER	166	9.832	12.064	-20.146	1.00	6.05	SEG1
ATOM	2567	N	GLY	167	9.947	10.605	-18.497	1.00	6.17	SEG1
ATOM	2568	HN	GLY	167	9.933	10.434	-17.533	1.00	6.15	SEG1
ATOM	2569	CA	GLY	167	10.115	9.466	-19.445	1.00	7.07	SEG1
ATOM	2570	HA1	GLY	167	9.356	9.521	-20.209	1.00	7.56	SEG1
ATOM	2571	HA2	GLY	167	10.016	8.534	-18.907	1.00	7.20	SEG1
ATOM	2572	C	GLY	167	11.500	9.536	-20.104	1.00	7.52	SEG1
ATOM	2573	O	GLY	167	11.679	9.107	-21.227	1.00	7.72	SEG1
ATOM	2574	N	ALA	168	12.485	10.078	-19.418	1.00	7.99	SEG1
ATOM	2575	HN	ALA	168	12.317	10.421	-18.514	1.00	8.05	SEG1
ATOM	2576	CA	ALA	168	13.867	10.183	-20.008	1.00	8.73	SEG1
ATOM	2577	HA	ALA	168	14.530	10.691	-19.325	1.00	8.71	SEG1
ATOM	2578	CB	ALA	168	14.342	8.737	-20.217	1.00	9.39	SEG1
ATOM	2579	HB1	ALA	168	13.732	8.068	-19.631	1.00	9.67	SEG1
ATOM	2580	HB2	ALA	168	15.373	8.647	-19.905	1.00	9.70	SEG1
ATOM	2581	HB3	ALA	168	14.258	8.477	-21.263	1.00	9.50	SEG1
ATOM	2582	C	ALA	168	13.814	10.921	-21.351	1.00	9.19	SEG1
ATOM	2583	O	ALA	168	12.802	11.496	-21.707	1.00	9.45	SEG1
ATOM	2584	N	ARG	169	14.889	10.898	-22.102	1.00	9.54	SEG1
ATOM	2585	HN	ARG	169	15.691	10.421	-21.799	1.00	9.50	SEG1
ATOM	2586	CA	ARG	169	14.885	11.589	-23.430	1.00	10.22	SEG1
ATOM	2587	HA	ARG	169	13.875	11.831	-23.723	1.00	10.32	SEG1
ATOM	2588	CB	ARG	169	15.717	12.879	-23.280	1.00	10.36	SEG1
ATOM	2589	HB1	ARG	169	15.480	13.541	-24.101	1.00	10.76	SEG1
ATOM	2590	HB2	ARG	169	16.766	12.635	-23.319	1.00	10.21	SEG1
ATOM	2591	CG	ARG	169	15.412	13.597	-21.959	1.00	10.51	SEG1
ATOM	2592	HG1	ARG	169	15.902	13.079	-21.148	1.00	10.29	SEG1
ATOM	2593	HG2	ARG	169	14.345	13.605	-21.789	1.00	10.79	SEG1
ATOM	2594	CD	ARG	169	15.931	15.037	-22.030	1.00	10.93	SEG1
ATOM	2595	HD1	ARG	169	15.512	15.627	-21.231	1.00	11.12	SEG1
ATOM	2596	HD2	ARG	169	15.682	15.474	-22.988	1.00	11.10	SEG1
ATOM	2597	NE	ARG	169	17.421	14.950	-21.875	1.00	11.16	SEG1
ATOM	2598	HE	ARG	169	17.987	14.946	-22.675	1.00	11.22	SEG1
ATOM	2599	CZ	APG	169	17.988	14.956	-20.684	1.00	11.49	SEG1
ATOM	2600	NH1	APG	169	19.290	15.019	-20.594	1.00	11.90	SEG1
ATOM	2601	HH11	APG	169	19.848	15.062	-21.421	1.00	11.96	SEG1
ATOM	2602	HH12	APG	169	19.728	15.028	-19.694	1.00	12.28	SEG1
ATOM	2603	NH2	APG	169	17.276	14.870	-19.586	1.00	11.63	SEG1
ATOM	2604	HH21	APG	169	16.282	14.794	-19.631	1.00	11.47	SEG1
ATOM	2605	HH22	APG	169	17.732	14.883	-18.697	1.00	12.03	SEG1

ATOM	2606	C	ARG	169	15.534	10.680	-24.477	1.00	10.92	SEG1
ATOM	2607	OT1	ARG	169	15.667	9.498	-24.206	1.00	11.30	SEG1
ATOM	2608	OT2	ARG	169	15.898	11.183	-25.529	1.00	11.23	SEG1
END										

FIG. 2 (35 of 35)

Relative binding affinities of TRADD-N mutants with TF2-C

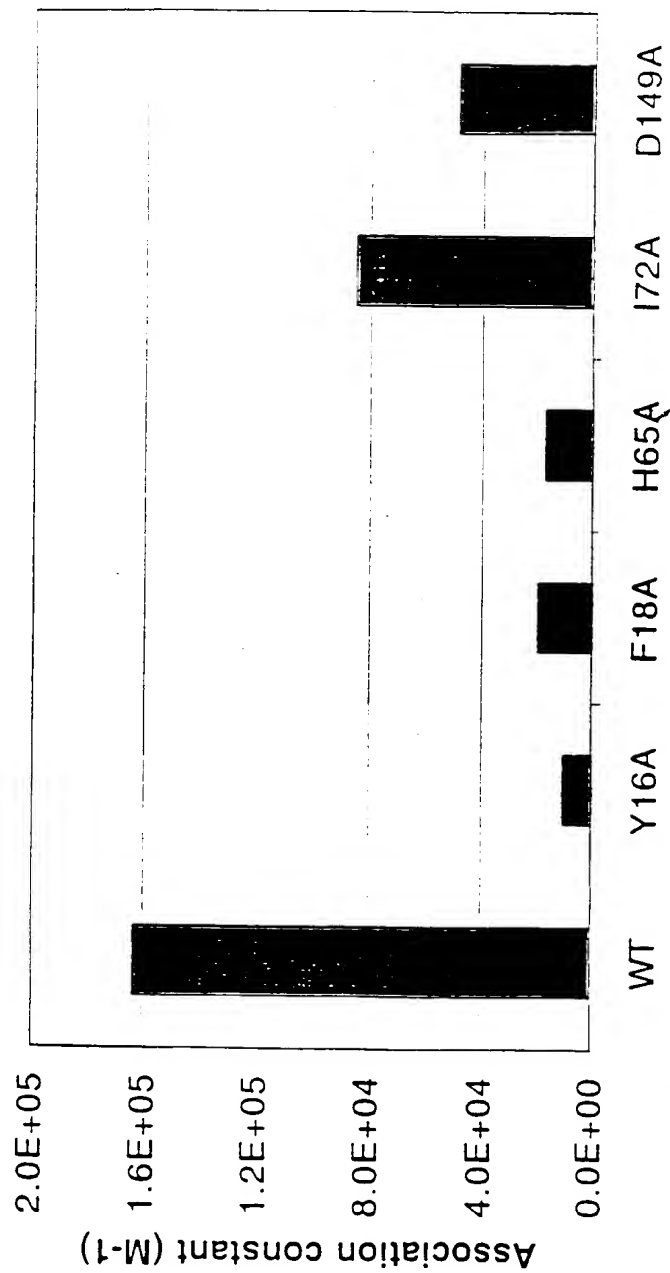


FIG. 3